



SEQUENCE LISTING

<110> Pramod K. Srivastava

<120> ALPHA(2) MACROGLOBULIN RECEPTOR AS A HEAT SHOCK
PROTEIN RECEPTOR AND USES THEREOF

<130> 8449-134

<140>

<141>

<150> 09/750,972

<151> 2000-12-28

<150> 09/668,724

<151> 2000-09-22

<160> 57

<170> FastSEQ for Windows Version 3.0

<210> 1

<211> 14849

<212> DNA

<213> Mus musculus

<400> 1

cgctgctccc	cgccagtgca	ctgaggaggc	ggaaacgggg	gagcccctag	tgctccatca	60
ggcccctacc	aaggcaccce	catcggggtcc	acgcccccca	ccccccaccc	cgctcctccc	120
caattgtgca	tttttgcagc	cggagtcggc	tccgagatgg	ggctgtgagc	ttcgccctgg	180
gagggggaga	ggagcgagga	gtaaagcagg	ggtgaagggt	tcgaatttgg	gggcaggggg	240
cgcaccgcgc	tcagcaggcc	cttcccaggg	ggctcggaac	tgtaaccattt	cacctatgcc	300
cctggttcgc	tttgcttaag	gaaggataag	atagaagagt	cggggagagg	aagataaagg	360
gggaccccc	aattgggggg	ggcgaggaca	agaagtaaca	ggaccagagg	gtgggggctg	420
ctgtttgcat	cggcccacac	catgctgacc	ccgccgttgc	tgctgctcgt	gccgctgctt	480
tcagctctgg	tctccggggc	cactatggat	gcccctaaaa	cttgcagccc	taagcagttt	540
gcctgcagag	accaaatac	ctgtatctca	aagggtctgg	ggtgtgacgg	tgaaagagat	600
tgccccgacg	gctctgatga	agcccctgag	atctgtccac	agagtaaagc	ccagagatgc	660
ccgccaaatg	agcacagttg	tctggggact	gagctatgtg	tccccatgtc	tcgtctctgc	720
aacgggateg	aggactgcat	ggatgggtca	gacgagggtg	ctcactgccg	agagctccga	780
gccaactgtt	ctcgaatggg	ttgtcaacac	cattgtgtac	ctacacccag	tgggcccacg	840
tgctactgta	acagcagctt	ccagctcgag	gcagatggca	agacgtgcaa	agattttgac	900
gagtgttccg	tgtatggcac	ctgcagccag	ctttgcacca	acacagatgg	ctccttcaca	960
tgtggctgtg	ttgaaggcta	cctgctgcaa	ccggacaacc	gctcctgcaa	ggccaagaat	1020
gagccagtag	atcgcccgcc	agtgtctactg	attgccaact	ctcagaacat	cctagctacg	1080
tacctgagtg	gggcccgaag	gtctaccatc	acaccaccca	gcacccgaca	aaccacggcc	1140
atggacttca	gttatgccaa	tgagaccgta	tgctgggtgc	acgttgggga	cagtgtgtcc	1200
cagacacagc	tcaagtgtgc	ccggatgcct	ggcctgaagg	gctttgtgga	tgagcatacc	1260
atcaacatct	ccctcagcct	gcaccacgtg	gagcagatgg	caatcgactg	gctgacggga	1320
aacttctact	ttgtcgacga	cattgacgac	aggatctttg	tctgtaaccg	aaacggggac	1380
acctgtgtca	ctctgtctga	cctggaactc	tacaacccca	aaggcatcgc	cttggacccc	1440
gccatgggga	aggtgttctt	cactgactac	gggcagatcc	caaagggtga	gcgctgtgac	1500
atggatggac	agaaccgcac	caagctgggtg	gatagcaaga	tcgtgtttcc	acacggcatc	1560
accctggacc	tggtcagccg	cctcgtctac	tgggcggacg	cctacctaga	ctacatcgag	1620
gtggtagact	acgaagggaa	gggtcggcag	accatcatcc	aaggcatcct	gatcgagcac	1680
ctgtacggcc	tgaccgtgtt	tgagaactat	ctctacgcca	ccaactcgga	caatgccaac	1740

acgcagcaga	agacgagcgt	gatccgagtg	aaccggttca	acagtactga	gtaccagggtc	1800
gtcaccgcgtg	tggacaaggg	tggtgccctg	catatctacc	accagcgacg	ccagcccccga	1860
gtgcggagtc	acgcctgtga	gaatgaccag	tacgggaagc	caggtggctg	ctccgacatc	1920
tgctcctgg	ccaacagtca	caaggcaagg	acctgcaggt	gcaggtctgg	cttcagcctg	1980
ggaagtgatg	ggaagtcttg	taagaaacct	gaacatgagc	tgttcctcgt	gtatggcaag	2040
ggccgaccag	gcatcattag	aggcatggac	atggggggcca	aggtcccaga	tgagcacatg	2100
atccccatcg	agaaccttat	gaatccacgc	gctctggact	tccacgccga	gaccggcttc	2160
atctactttg	ctgacaccac	cagctacctc	attggccgcc	agaaaattga	tggcacggag	2220
agagagacta	tcctgaagga	tggcatccac	aatgtggagg	gcgtagccgt	ggactggatg	2280
ggagacaatc	tttactggac	tgatgatggc	cccaagaaga	ccattagtgt	ggccaggctg	2340
gagaaagccg	ctcagacccg	gaagactcta	attgagggca	agatgacaca	ccccagggcc	2400
attgtagtgg	atccactcaa	tgggtggatg	tactggacag	actgggagga	ggaccccaag	2460
gacagtcggc	gagggcggct	cgagagggct	tggatggacg	gctcacaccg	agatatcttt	2520
gtcacctcca	agacagtgct	ttggcccaat	gggctaagcc	tggatatccc	agccggacgc	2580
ctctactggg	tggatgcctt	ctatgaccga	attgagacca	tactgctcaa	tggcacagac	2640
cggaaagattg	tatatgaggg	tcctgaactg	aatcatgcct	tcggcctgtg	tcaccatggc	2700
aactacctct	tttggaccga	gtaccggagc	ggcagcgtct	accgcttgga	acggggcgctg	2760
gcaggcgcac	cgcccaactgt	gacccttctg	cgcagcgaga	gaccgcctat	ctttgagatc	2820
cgaatgtacg	acgcgcacga	gcagcaagtg	ggtaccaaca	aatgccgggt	aaataacgga	2880
ggctgcagca	gcctgtgcct	cgccaccccc	gggagccgcc	agtgtgcctg	tgccgaggac	2940
caggtgttgg	acacagatgg	tgtcacctgc	ttggcgaacc	catcctacgt	gccccacccc	3000
cagtgccagc	cgggccagtt	tgctgtgcc	aacaaccgct	gcatccagga	gcgctggaag	3060
tgtgacggag	acaacgactg	tctggacaac	agcgatgagg	cccagcact	gtgccatcaa	3120
cacacctgtc	cctcggaccg	attcaagtgt	gagaacaacc	ggtgtatccc	caaccgctgg	3180
ctctgtgatg	gggataatga	ttgtggcaac	agcgaggacg	aatccaatgc	cacgtgctca	3240
gcccgcacct	gtccacccaa	ccagttctcc	tgtgccagtg	gccgatgcat	tcctatctca	3300
tggacctgtg	atctggatga	tgactgtggg	gaccggtccg	atgagtcagc	ctcatgcgcc	3360
taccccaacct	gcttccccct	gactcaattt	acctgcaaca	atggcagatg	tattaacatc	3420
aactggcgggt	gtgacaacga	caatgactgt	ggggacaaca	gcgacgaagc	cggctgcagt	3480
cactcctgct	ccagtaccca	gttcaagtgc	aacagtggca	gatgcatccc	cgagcactgg	3540
acgtgtgatg	gggacaatga	ttgtggggac	tacagcgacg	agacacacgc	caactgtacc	3600
aaacaggcta	caagacctcc	tgggtggctgc	cactcggatg	agttccagtg	cccgctagat	3660
ggcctgtgca	tccccctgag	gtggcgctgc	gacggggaca	ccgactgcat	ggattccagc	3720
gatgagaaga	gctgtgaggg	cgtgacctat	gtttgtgacc	cgaatgtcaa	gtttggctgc	3780
aaggactccg	cccggtgcat	cagcaaggcg	tgggtgtgtg	atggcgacag	cgactgtgaa	3840
gataactccg	acgaggagaa	ctgtgaggcc	ctggcctgca	ggccaccctc	ccatccctgc	3900
gccaacaaca	cctctgtctg	cctgcctcct	gacaagctgt	gcgacggcaa	ggatgactgt	3960
ggagacggct	cggatgaggg	cgagctctgt	gaccagtgtt	ctctgaataa	tgggtggctgt	4020
agtcacaact	gctcagtggc	ccctggtgaa	ggcatcgtgt	gctcttgccc	tctgggcatg	4080
gagctgggct	ctgacaacca	cacctgccag	atccagagct	actgtgccaa	gcacctcaaa	4140
tgacagccaga	agtgtgacca	gaacaagttc	agtgtgaagt	gctcctgcta	cgagggctgg	4200
gtcttggagc	ctgacgggga	aacgtgccgc	agtctggatc	ccttcaaact	gttcatcatc	4260
ttctccaacc	gccacgagat	caggcgcatt	gaccttcaca	agggggacta	cagcgtccta	4320
gtgcctggcc	tgcgcaacac	tattgccttg	gacttccacc	tcagccagag	tgccctctac	4380
tggaccgacg	cggtagagga	caagatctac	cgtgggaaac	tcctggacaa	cggagccctg	4440
accagctttg	aggtgggtgat	tcagtatggc	ttggccacac	cagagggcct	ggctgtagat	4500
tggattgcag	gcaacatcta	ctgggtggag	agcaacctgg	accagatcga	agtggccaag	4560
ctggacggaa	ccctccgaac	cactctgctg	gcgggtgaca	ttgagcacc	gagggccatc	4620
gctctggacc	ctcgggatgg	gattctgttt	tggacagact	gggatgccag	cctgccacga	4680
atcgaggctg	catccatgag	tggagctggc	cgccgaacca	tccaccggga	gacaggctct	4740
gggggctgcg	ccaatgggct	caccgtggat	tacctggaga	agcgcatact	ctggattgat	4800
gctaggtcag	atgccatcta	ttcagcccgg	tatgacggct	ccggccacat	ggaggtgctt	4860
cggggacacg	agttcctgtc	acacccattt	gccgtgacac	tgtacggtgg	ggaggtgtac	4920
tggaccgact	ggcgaacaaa	tacactggct	aaggccaaca	agtggactgg	ccacaacgtc	4980
accgtggtac	agaggaccaa	caccagccc	ttcgacctgc	aggtgtatca	cccttcccgg	5040
cagcccatgg	ctccaaaccc	atgtgaggcc	aatggcggcc	ggggcccctg	ttcccatctg	5100
tgctcatca	actacaaccg	gaccgtctcc	tgggcctgtc	cccacctcat	gaagctgcac	5160
aaggacaaca	ccacctgcta					

atccggggcg	tggacctgga	tgccccgtac	tacaattata	tcattctcctt	cacgggtgcct	5280
gatatcgaca	atgtcacggg	gctggactat	gatgcccag	agcagcgagt	ttactgggtct	5340
gatgtgcgga	ctcaagccat	caaaagggca	tttatcaacg	gcactggcgt	ggagaccgtt	5400
gtctctgcag	acttgcccaa	cgcccacggg	ctggctgtgg	actgggtctc	ccgaaatctg	5460
ttttggacaa	gttacgacac	caacaagaag	cagattaacg	tggcccggct	ggacggctcc	5520
ttcaagaatg	cgggtgggtgca	gggcctggag	cagccccacg	gcctgggtcgt	ccacccgctt	5580
cgtggcaagc	tctactggac	tgatggggac	aacatcagca	tggccaacat	ggatggggagc	5640
aaccacactc	tgctcttcag	tggccagaag	ggccctgtgg	ggttggccat	tgacttcctt	5700
gagagcaaac	tctactggat	cagctctggg	aaccacacaa	tcaaccgttg	caatctggat	5760
gggagcgagc	tggaggtcat	cgacaccatg	cggagccagc	tgggcaaggc	cactgccctg	5820
gccatcatgg	gggacaagct	gtgggtgggca	gatcaggtgt	cagagaagat	gggcacgtgc	5880
aacaaagccg	atggctctgg	gtccgtgggtg	ctgcggaaca	gtaccacgtt	ggttatgcac	5940
atgaaggtgt	atgacgagag	catccagcta	gagcatgagg	gcaccaaccc	ctgcagtgtc	6000
aacaacggag	actgttccca	gctctgcctg	ccaacatcag	agacgactcg	ctcctgtatg	6060
tgtacagccg	gttacagcct	ccggagcgga	cagcaggcct	gtgaggggtgt	gggctctttt	6120
ctcctgtact	ctgtacatga	gggaattcgg	gggattccac	tagatcccaa	tgacaagtcg	6180
gatgccctgg	tcccagtgct	cggaaacttca	ctggctgtcg	gaatcgactt	ccatgccgaa	6240
aatgacacta	tttattgggt	ggatatgggc	ctaagcacca	tcagcagggc	caagcgtgac	6300
cagacatggc	gagaggatgt	ggtgaccaac	ggtattggcc	gtgtggaggg	catcgccgtg	6360
gactggatcg	caggcaacat	atactggacg	gaccagggct	tcgatgtcat	cgaggttgcc	6420
cggctcaatg	gctcttttcg	ttatgtgggtc	atttcccagg	gtctggacaa	gcctcggggc	6480
atcactgtcc	accagagaaa	ggggactttg	ttctggaccg	agtgggggtca	ttacccacgt	6540
attgagcggg	ctcgccctga	tggcacagag	agagtgggtgt	tggttaatgt	cagcatcagc	6600
tggcccaatg	gcattctcagt	agactatcag	ggcggcaagc	tctactgggtg	tgatgctcgg	6660
atggacaaga	tcgagcgcct	cgacctggaa	acggggcgaga	accggggaggt	ggtcctgtcc	6720
agcaataaca	tggatatgtt	ctccgtgtcc	gtgtttgagg	acttcatcta	ctggagtgc	6780
agaactcacg	ccaatggctc	catcaagcgc	ggctgcaaag	acaatgctac	agactccgtg	6840
cctctgagga	caggcattgg	tgttcagctt	aaagacatca	aggtcttcaa	cagggacagg	6900
cagaagggta	ccaatgtgtg	cgcggtagcc	aacggcgggg	gccagcagct	ctgcttgtat	6960
cgggggtggcg	gacagcgagc	ctgtgcctgt	gccacgggga	tgctggcaga	agacgggggc	7020
tcattgccgag	agtagcgtgg	ctacctgtct	tactcagagc	ggaccatcct	caagagcatc	7080
cacctgtcgg	atgagcgtaa	cctcaacgca	ccggtgcagc	cctttgaaga	ccccgagcac	7140
atgaaaaatg	tcattgcctt	ggcctttgac	taccgagcag	gcacctcccc	ggggaccctt	7200
aaccgcatct	tcttcagtga	catccacttt	gggaacatcc	agcagatcaa	tgacgatggc	7260
tcgggcagga	ccaccatcgt	ggaaaatgtg	ggctctgtgg	aaggcctggc	ctatcacctg	7320
ggctgggaca	cactgtactg	gacaagctac	accacatcca	ccatcacccg	ccacaccgtg	7380
gaccagactc	gccagggggc	cttcgagagg	gagacagtca	tcaccatgtc	cggagacgac	7440
cacccgagag	cctttgtgct	ggatgagtgc	cagaacctga	tgttctggac	caattggaac	7500
gagctccatc	caagcatcat	gcgggcagcc	ctatccggag	ccaacgtcct	gaccctcatt	7560
gagaaggaca	tccgcacgcc	caatgggttg	gccatcgacc	accggggcgga	gaagctgtac	7620
ttctcggatg	ccaccttgga	caagatcgag	cgtgcgagt	acgacggctc	ccaccgctat	7680
gtgatcctaa	agtcggagcc	cgtccacccc	tttgggttgg	cgggtgtacgg	agagcacatt	7740
ttctggactg	actgggtgcg	gcgggctgtg	cagcgagcca	acaagtatgt	gggcagcgac	7800
atgaagctgc	ttcgggtgga	cattccccag	caacccatgg	gcattcatcg	cgtggccaat	7860
gacaccaaca	gctgtgaact	ctccccctgc	cgtatcaaca	atggaggctg	ccaggatctg	7920
tgtctgctca	cccaccaagg	ccacgtcaac	tgttctgtgc	gagggggccg	gattctccag	7980
gaggacttca	cctgccgggc	tgtgaactcc	tcttgtcggg	cacaagatga	gtttgagtgt	8040
gccaatgggg	aatgtatcag	cttcagcctc	acctgtgatg	gcgtctccca	ctgcaaggac	8100
aagtccgatg	agaagccctc	ctactgcaac	tcacgccgct	gcaagaagac	tttccgcccag	8160
tgtaacaatg	gccgctgtgt	atccaacatg	ctgtgggtgca	atgggggtgga	ttactgtggg	8220
gatggctctg	atgagatacc	ttgcaacaag	actgctgtg	gtgtgggtga	gttccgctgc	8280
cgggatgggt	cctgcacgag	gaactccagt	cgtgcgaacc	agtttgtgga	ttgtgaggat	8340
gcctcggatg	agatgaattg	cagtgccaca	gactgcagca	gctatttccg	cctgggcgtg	8400
aaaggtgtcc	tcttccagcc	gtgcgagcgg	acatccctgt	gctacgcacc	tagctgggtg	8460
tgtgatggcg	ccaacgactg	tggagactac	agcgatgaac	gtgactgtcc	aggtgtgaag	8520
cgccttaggt	gcccgtcaa	ttactttgcc	tgccccagcg	ggcgtgtgat	ccccatgagc	8580
tggacgtgtg	acaaggagga	tgactgtgag	aacggcgagg	atgagaccca	ctgcaacaag	8640
ttctgctcag	aggcacagtt	cgagtgccag	aaccaccggg	gtattctcaa	gcagtggctg	8700

tgtgacggta	gcgatgattg	cggggatggc	tccgatgagg	cagctcactg	tgaaggcaag	8760
acatgtggcc	cctcctcctt	ctcctgtccc	ggcaccacg	tgtgtgtccc	tgagcgctgg	8820
ctctgtgatg	gcgacaagga	ctgtaccgat	ggcgcggatg	agagtgtcac	tgctggctgc	8880
ctgtacaaca	gcacctgtga	tgaccgtgag	ttcatgtgcc	agaaccgctt	gtgtattccc	8940
aagcatttcg	tgtgcgacca	tgaccgtgac	tgtgctgatg	gctctgatga	atccccctgag	9000
tgtgagtacc	caacctgcgg	gcccgaatgaa	ttccgctgtg	ccaatggggcg	ttgtctgagc	9060
tcccgtcagt	gggaatgtga	tggggagaaat	gactgtcacg	accacagcga	tgaggctccc	9120
aagaaccac	actgcaccag	cccagagcac	aaatgcaatg	cctcatcaca	gttcctgtgc	9180
agcagcgggc	gctgcgtggc	tgaggcggtg	ctctgcaacg	gccaggacga	ctgtggggac	9240
ggttcagacg	aacgcgggtg	ccatgtcaac	gagtgtctca	gccgcaagct	cagtggctgc	9300
agtcaggact	gcgaggacct	caagataggc	tttaagtgcc	gctgtcgccc	gggcttccgg	9360
ctaaaggacg	atggcaggac	ctgtgccgac	ctggatgagt	gcagcaccac	cttccccctgc	9420
agccagctct	gcatcaacac	ccacggaagt	tacaagtgtc	tgtgtgtgga	gggctatgca	9480
ccccgtggcg	gtgaccccca	cagctgcaaa	gctgtgaccg	atgaggagcc	atttctcatc	9540
tttgccaacc	ggtactacct	gcggaagctc	aacctggacg	gctccaacta	cacactgctt	9600
aagcagggcc	tgaacaatgc	ggtcgccttg	gcatttgact	accgagagca	gatgatctac	9660
tggacggggcg	tgaccacca	gggcagcatg	attcgcagga	tgcacctcaa	cggcagcaac	9720
gtgcaggttc	tgcaccggac	gggccttagt	aaccagatg	ggctcgctgt	ggactgggtg	9780
ggtggcaacc	tgtactgggtg	tgacaagggc	agagatacca	ttgagggtgtc	caagcttaac	9840
ggggcctatc	ggacagtgtc	ggtcagctct	ggcctccggg	agcccagagc	tctggtagtg	9900
gatgtacaga	atgggtacct	gtactggaca	gactgggggtg	accactcact	gatcggccgg	9960
attggcatgg	atggatctgg	ccgcagcatc	atcgtggaca	ctaagatcac	atggcccaat	10020
ggcctgaccg	tggactacgt	cacggaacgc	atctactggg	ctgacgcccg	tgaggactac	10080
atcgagttcg	ccagcctgga	tggctccaac	cgtcacgttg	tgctgagcca	agacatccca	10140
cacatctttg	cgctgaccct	atttgaagac	tacgtctact	ggacagactg	ggaaacgaag	10200
tccatcaacc	gggcccacaa	gaccacgggt	gccaaacaaa	cactcctcat	cagcacccctg	10260
caccggccca	tggacttaca	tgtattccac	gccctgcgcc	agccagatgt	gcccgaatcac	10320
ccctgcaaag	tcaacaatgg	tggctgcagc	aacctgtgcc	tgctgtcccc	tgggggtggt	10380
cacaagtgcg	cctgccccac	caacttctat	ctgggtggcg	atggccgtac	ctgtgtgtcc	10440
aactgcacag	caagccagtt	tgtgtgcaaa	aatgacaagt	gcatccccctt	ctggtggaag	10500
tgtgacacgg	aggacgactg	tggggatcac	tcagacgagc	ctccagactg	tcccgagttc	10560
aagtgccgcc	caggccagtt	ccagtgtctc	accggcatct	gcaccaaccc	tgccttcctc	10620
tgtgatgggg	acaatgactg	ccaagacaat	agtgacgagg	ccaattgcga	cattcacgtc	10680
tgcttgccca	gccaattcaa	gtgcaccaac	accaaccgct	gcattcctgg	catcttccgt	10740
tgcaatgggc	aggacaactg	cggggacggc	gaggatgagc	gggattgccc	tgaggtgacc	10800
tgcgccccca	accagttcca	gtgctccatc	accaagcgct	gcatccctcg	cgtctgggtc	10860
tgtgacaggg	ataatcactg	tgtggacggc	agtgatgagc	ctgccaaactg	tacccaaatg	10920
acctgtggag	tggatgagtt	ccgctgcaag	gattctggcc	gctgcatccc	cgcgcgctgg	10980
aagtgtgacg	gagaagatga	ctgtggggat	ggttcagatg	agcccaagga	agagtgtgat	11040
gagcgcacct	gtgagccata	ccagttccgc	tgcaaaaaca	accgctgtgt	cccaggccgt	11100
tggcaatgtg	actacgacaa	cgactgcgga	gataactcgg	acgaggagag	ctgcacacct	11160
cggccctgct	ctgagagtga	gtttttctgt	gccaatggcc	gctgcatcgc	tgggcgctgg	11220
aagtgtgatg	gggaccatga	ctgtgccgac	ggctcagacg	agaaagactg	caccccccg	11280
tgtgatatgg	accagttcca	gtgcaagagt	ggccactgca	tccccctgcg	ctggccgtgt	11340
gacgcggatg	ctgactgtat	ggacggcagt	gacgaggaag	cctgtggcac	tggggtgagg	11400
acctgcccat	tggatgagtt	tcaatgtaac	aacaccttgt	gcaagccgct	ggcctggaag	11460
tgtgatggag	aggacgactg	tggggacaac	tcagatgaga	accccagagga	atgcgcccgg	11520
ttcatctgcc	ctcccaaccg	gcctttccgc	tgcaagaatg	accgagtctg	cctgtggatt	11580
gggcgccagt	gtgatggcgt	ggacaactgt	ggagatggga	ctgacgagga	ggactgtgag	11640
ccccccacgg	cccagaaccc	ccactgcaaa	gacaagaagg	agttcctgtg	ccgaaaccag	11700
cgctgtctat	catcctccct	gcgctgtaac	atgttcgatg	actgcggcga	tggctccgat	11760
gaagaagatt	gcagcatcga	ccccaaagctg	accagctgtg	ccaccaatgc	cagcatgtgt	11820
ggggacgaag	ctcgttgtgt	gcgcactgag	aaagctgcct	actgtgcctg	ccgctcgggc	11880
ttccatactg	tgccggggcca	gcccgggatgc	caggacatca	acgagtgcct	gcgctttggt	11940
acctgctctc	agctctggaa	caaaccacaag	ggaggccacc	tctgcagctg	tgcccgcac	12000
ttcatgaaga	cacacaacac	ctgcaaagct	gaaggctccg	agtaccaggt	gctatacatc	12060
gcggatgaca	acgagatccg	cagcttgttc	ccgggccacc	cccactcagc	ctacg	

cgtgtctact	ggactaactg	gcacacgggc	acaatctcct	acaggagcct	gccccctgcc	12240
gccccctcta	ccacttccaa	ccgccaccgg	aggcagatcg	accggggtgt	caccacacctc	12300
aatatttccag	ggctgaagat	gccgaggggt	atcgctatcg	actgggtggc	cgggaatgtg	12360
tactggaccg	attccggccg	agacgtgatt	gaggtggcgc	aaatgaaggg	cgagaaccgc	12420
aagacgctca	tctcgggcat	gattgatgag	ccccatgcca	tcgtgggtgga	ccctctgagg	12480
ggcaccatgt	actggtcaga	ctgggggaac	cacccaaga	ttgaaacagc	agcgatggat	12540
ggcacccttc	gggagactct	cgtgcaagac	aacattcagt	ggcctacagg	gctggctgtg	12600
gactatcaca	atgaacggct	ctactgggca	gatgccaagc	tttcgggtcat	cggcagcatc	12660
cggctcaacg	gcactgaccc	cattgtggct	gctgacagca	aacgaggcct	aagtcacccc	12720
ttcagcatcg	atgtgtttga	agactacatc	tacggagtca	cttacatcaa	taatcgtgtc	12780
ttcaagatcc	acaagtttgg	acacagcccc	ttgtacaacc	taactggggg	cctgagccat	12840
gcctctgatg	tagtccttta	ccatcaacac	aagcagcctg	aagtgaccaa	cccctgtgac	12900
cgcaagaaat	gcgaatggct	gtgtctgctg	agccccagcg	ggcctgtctg	cacctgtccc	12960
aatggaaaga	ggctggataa	tggcacctgt	gtgcctgtgc	cctctccaac	acccccctca	13020
gatgccccta	ggcctggaac	ctgcactctg	cagtgcctca	atgggtggtag	ttgtttcctc	13080
aacgctcgga	ggcagcccaa	gtgccgttgc	cagccccgtt	acacaggcga	taagtgtgag	13140
ctggatcagt	gctgggaata	ctgtcacaa	ggaggcacct	gtgcggcttc	cccatctggc	13200
atgcccacgt	gccgctgtcc	cactggcttc	acgggccccca	aatgcaccgc	acaggtgtgt	13260
gcaggctact	gctctaacaa	cagcacctgc	accgtcaacc	agggcaacca	gccccagtgc	13320
cgatgtctac	ctggcttcct	gggcgaccgt	tgccagtacc	ggcagtgtct	tggcttctgt	13380
gagaactttg	gcacctgtca	gatggctgct	gatggctccc	gacaatgtcg	ctgcaccgtc	13440
tactttgagg	gaccaagggtg	tgagggtgaac	aagtgtagtc	gctgtctcca	aggcgctgt	13500
gtggtcaata	agcagaccgg	agatgtcaca	tgcaactgca	ctgatggccg	ggtagcccc	13560
agttgtctca	cctgcacoga	tcactgtagc	aatgggtggct	cctgcaccat	gaacagcaag	13620
atgatgcctg	agtgccagtg	cccgccccat	atgacaggac	cccggtgcca	ggagcagggtt	13680
gttagtcagc	aacagcctgg	gcatatggcc	tccatcctga	tccctctgct	gctgcttctc	13740
ctgctgcttc	tgggtggctgg	cgtgggtgttc	tggtataaagc	ggcgagtccg	aggggctaag	13800
ggcttccagc	accagcggat	gaccaatggg	gccatgaatg	tggaaattgg	aaaccctacc	13860
tacaagatgt	atgaagggtgg	agagcccgat	gatgtcgggg	gcctactgga	tgctgatttt	13920
gcccttgacc	ctgacaagcc	taccaacttc	accaaccag	tgtatgccac	gctctacatg	13980
ggggggccacg	gcagccgcca	ttccctggcc	agcacggacg	agaagcgaga	actgctgggc	14040
cggggacctg	aagacgagat	aggagatccc	ttggcatagg	gccctgcccc	gacggatgtc	14100
cccagaaaagc	cccctgccac	atgagtcttt	caatgaaccc	cctccccagc	cggcccttct	14160
ccggccctgc	cgggtgtaca	aatgtaaaaa	tgaaggaatt	actttttata	tgtgagcgag	14220
caagcgagca	agcacagtat	tatctctttg	catttccttc	ctgcctgtct	ctcagtatcc	14280
cccccatgct	gccttgaggg	ggcggggagg	gctttgtggc	tcaaagggtat	gaaggagtcc	14340
acatgttccc	taccgagcat	acccttgga	gcctggcggc	acggcctccc	caccacgcct	14400
gtgcaagaca	ctcaacgggg	ctccgtgtcc	cagctttcct	ttccttggct	ctctgggggtt	14460
agttcagggg	aggtggagtc	ctctgctgac	cctgtctgga	agatttggct	ctagctgagg	14520
aaggagtctt	ttagttaggg	gaagtcaccc	caaaccaccag	ctcccacttt	caggggcacc	14580
tctcagatgg	ccatgctcag	tatcccttcc	agacaggccc	tccctctctt	agcgccccct	14640
ctgtggctcc	tagggctgaa	cacattcttt	ggtaactgtc	ccccaagcct	cccatcccc	14700
tgagggccag	gaagagtcgg	ggcacaccaa	ggaagggcaa	gcgggcagcc	ccatttttggg	14760
gacgtgaacg	ttttaataat	ttttgctgaa	ttcctttaca	actaaataac	acagatattg	14820
ttataaataa	aatttqtaaaa	aaaaaaaaa				14849

```
<210> 2
<211> 4545
<212> PRT
<213> Mus musculus
```

```

      <400> 2
Met  Leu  Thr  Pro  Pro  Leu  Leu  Leu  Leu  Val  Pro  Leu  Leu  Ser  Ala  Leu
 1          5          10          15
Val  Ser  Gly  Ala  Thr  Met  Asp  Ala  Pro  Lys  Thr  Cys  Ser  Pro  Lys  Gln
      20          25          30

```


Ala	Asn	Ser	His	Lys	Ala	Arg	Thr	Cys	Arg	Cys	Arg	Ser	Gly	Phe	Ser	
			500					505					510			
Leu	Gly	Ser	Asp	Gly	Lys	Ser	Cys	Lys	Lys	Pro	Glu	His	Glu	Leu	Phe	
		515					520					525				
Leu	Val	Tyr	Gly	Lys	Gly	Arg	Pro	Gly	Ile	Ile	Arg	Gly	Met	Asp	Met	
	530					535					540					
Gly	Ala	Lys	Val	Pro	Asp	Glu	His	Met	Ile	Pro	Ile	Glu	Asn	Leu	Met	
545					550					555					560	
Asn	Pro	Arg	Ala	Leu	Asp	Phe	His	Ala	Glu	Thr	Gly	Phe	Ile	Tyr	Phe	
			565					570						575		
Ala	Asp	Thr	Thr	Ser	Tyr	Leu	Ile	Gly	Arg	Gln	Lys	Ile	Asp	Gly	Thr	
			580					585					590			
Glu	Arg	Glu	Thr	Ile	Leu	Lys	Asp	Gly	Ile	His	Asn	Val	Glu	Gly	Val	
	595					600						605				
Ala	Val	Asp	Trp	Met	Gly	Asp	Asn	Leu	Tyr	Trp	Thr	Asp	Asp	Gly	Pro	
	610				615						620					
Lys	Lys	Thr	Ile	Ser	Val	Ala	Arg	Leu	Glu	Lys	Ala	Ala	Gln	Thr	Arg	
625					630					635					640	
Lys	Thr	Leu	Ile	Glu	Gly	Lys	Met	Thr	His	Pro	Arg	Ala	Ile	Val	Val	
			645					650						655		
Asp	Pro	Leu	Asn	Gly	Trp	Met	Tyr	Trp	Thr	Asp	Trp	Glu	Glu	Asp	Pro	
		660						665					670			
Lys	Asp	Ser	Arg	Arg	Gly	Arg	Leu	Glu	Arg	Ala	Trp	Met	Asp	Gly	Ser	
	675					680						685				
His	Arg	Asp	Ile	Phe	Val	Thr	Ser	Lys	Thr	Val	Leu	Trp	Pro	Asn	Gly	
	690					695					700					
Leu	Ser	Leu	Asp	Ile	Pro	Ala	Gly	Arg	Leu	Tyr	Trp	Val	Asp	Ala	Phe	
705					710					715					720	
Tyr	Asp	Arg	Ile	Glu	Thr	Ile	Leu	Leu	Asn	Gly	Thr	Asp	Arg	Lys	Ile	
			725					730						735		
Val	Tyr	Glu	Gly	Pro	Glu	Leu	Asn	His	Ala	Phe	Gly	Leu	Cys	His	His	
			740					745					750			
Gly	Asn	Tyr	Leu	Phe	Trp	Thr	Glu	Tyr	Arg	Ser	Gly	Ser	Val	Tyr	Arg	
	755					760						765				
Leu	Glu	Arg	Gly	Val	Ala	Gly	Ala	Pro	Pro	Thr	Val	Thr	Leu	Leu	Arg	
	770					775						780				
Ser	Glu	Arg	Pro	Pro	Ile	Phe	Glu	Ile	Arg	Met	Tyr	Asp	Ala	His	Glu	
785					790					795					800	
Gln	Gln	Val	Gly	Thr	Asn	Lys	Cys	Arg	Val	Asn	Asn	Gly	Gly	Cys	Ser	
			805					810						815		
Ser	Leu	Cys	Leu	Ala	Thr	Pro	Gly	Ser	Arg	Gln	Cys	Ala	Cys	Ala	Glu	
		820						825					830			
Asp	Gln	Val	Leu	Asp	Thr	Asp	Gly	Val	Thr	Cys	Leu	Ala	Asn	Pro	Ser	
	835					840						845				
Tyr	Val	Pro	Pro	Pro	Gln	Cys	Gln	Pro	Gly	Gln	Phe	Ala	Cys	Ala	Asn	
	850					855					860					
Asn	Arg	Cys	Ile	Gln	Glu	Arg	Trp	Lys	Cys	Asp	Gly	Asp	Asn	Asp	Cys	
865					870					875					880	
Leu	Asp	Asn	Ser	Asp	Glu	Ala	Pro	Ala	Leu	Cys	His	Gln	His	Thr	Cys	
			885					890						895		
Pro	Ser	Asp	Arg	Phe	Lys	Cys	Glu	Asn	Asn	Arg	Cys	Ile	Pro	Asn	Arg	
		900						905					910			
Trp	Leu	Cys	Asp	Gly	Asp	Asn	Asp	Cys	Gly	Asn	Ser	Glu	Asp	Glu	Ser	
	915					920						925				
Asn	Ala	Thr	Cys	Ser	Ala	Arg	Thr	Cys	Pro	Pro	Asn	Gln	Phe	Ser	Cys	
	930					935					940					
Ala	Ser	Gly	Arg	Cys	Ile	Pro	Ile	Ser	Trp	Thr	Cys	Asp	Leu	Asp	Asp	
945					950					955					960	

Asp	Cys	Gly	Asp	Arg	Ser	Asp	Glu	Ser	Ala	Ser	Cys	Ala	Tyr	Pro	Thr		
				965					970					975			
Cys	Phe	Pro	Leu	Thr	Gln	Phe	Thr	Cys	Asn	Asn	Gly	Arg	Cys	Ile	Asn		
			980					985					990				
Ile	Asn	Trp	Arg	Cys	Asp	Asn	Asp	Asn	Asp	Cys	Gly	Asp	Asn	Ser	Asp		
		995					1000					1005					
Glu	Ala	Gly	Cys	Ser	His	Ser	Cys	Ser	Ser	Thr	Gln	Phe	Lys	Cys	Asn		
	1010					1015					1020						
Ser	Gly	Arg	Cys	Ile	Pro	Glu	His	Trp	Thr	Cys	Asp	Gly	Asp	Asn	Asp		
1025					1030					1035					1040		
Cys	Gly	Asp	Tyr	Ser	Asp	Glu	Thr	His	Ala	Asn	Cys	Thr	Asn	Gln	Ala		
				1045					1050					1055			
Thr	Arg	Pro	Pro	Gly	Gly	Cys	His	Ser	Asp	Glu	Phe	Gln	Cys	Pro	Leu		
			1060					1065					1070				
Asp	Gly	Leu	Cys	Ile	Pro	Leu	Arg	Trp	Arg	Cys	Asp	Gly	Asp	Thr	Asp		
	1075					1080					1085						
Cys	Met	Asp	Ser	Ser	Asp	Glu	Lys	Ser	Cys	Glu	Gly	Val	Thr	His	Val		
	1090					1095					1100						
Cys	Asp	Pro	Asn	Val	Lys	Phe	Gly	Cys	Lys	Asp	Ser	Ala	Arg	Cys	Ile		
1105					1110					1115					1120		
Ser	Lys	Ala	Trp	Val	Cys	Asp	Gly	Asp	Ser	Asp	Cys	Glu	Asp	Asn	Ser		
				1125					1130					1135			
Asp	Glu	Glu	Asn	Cys	Glu	Ala	Leu	Ala	Cys	Arg	Pro	Pro	Ser	His	Pro		
			1140					1145					1150				
Cys	Ala	Asn	Asn	Thr	Ser	Val	Cys	Leu	Pro	Pro	Asp	Lys	Leu	Cys	Asp		
	1155						1160					1165					
Gly	Lys	Asp	Asp	Cys	Gly	Asp	Gly	Ser	Asp	Glu	Gly	Glu	Leu	Cys	Asp		
	1170					1175					1180						
Gln	Cys	Ser	Leu	Asn	Asn	Gly	Gly	Cys	Ser	His	Asn	Cys	Ser	Val	Ala		
1185					1190					1195					1200		
Pro	Gly	Glu	Gly	Ile	Val	Cys	Ser	Cys	Pro	Leu	Gly	Met	Glu	Leu	Gly		
				1205					1210					1215			
Ser	Asp	Asn	His	Thr	Cys	Gln	Ile	Gln	Ser	Tyr	Cys	Ala	Lys	His	Leu		
		1220						1225					1230				
Lys	Cys	Ser	Gln	Lys	Cys	Asp	Gln	Asn	Lys	Phe	Ser	Val	Lys	Cys	Ser		
	1235						1240					1245					
Cys	Tyr	Glu	Gly	Trp	Val	Leu	Glu	Pro	Asp	Gly	Glu	Thr	Cys	Arg	Ser		
	1250					1255					1260						
Leu	Asp	Pro	Phe	Lys	Leu	Phe	Ile	Ile	Phe	Ser	Asn	Arg	His	Glu	Ile		
1265					1270				1275						1280		
Arg	Arg	Ile	Asp	Leu	His	Lys	Gly	Asp	Tyr	Ser	Val	Leu	Val	Pro	Gly		
				1285					1290					1295			
Leu	Arg	Asn	Thr	Ile	Ala	Leu	Asp	Phe	His	Leu	Ser	Gln	Ser	Ala	Leu		
		1300						1305					1310				
Tyr	Trp	Thr	Asp	Ala	Val	Glu	Asp	Lys	Ile	Tyr	Arg	Gly	Lys	Leu	Leu		
	1315						1320					1325					
Asp	Asn	Gly	Ala	Leu	Thr	Ser	Phe	Glu	Val	Val	Ile	Gln	Tyr	Gly	Leu		
	1330					1335					1340						
Ala	Thr	Pro	Glu	Gly	Leu	Ala	Val	Asp	Trp	Ile	Ala	Gly	Asn	Ile	Tyr		
1345					1350				1355						1360		
Trp	Val	Glu	Ser	Asn	Leu	Asp	Gln	Ile	Glu	Val	Ala	Lys	Leu	Asp	Gly		
				1365					1370					1375			
Thr	Leu	Arg	Thr	Thr	Leu	Leu	Ala	Gly	Asp	Ile	Glu	His	Pro	Arg	Ala		
			1380					1385					1390				
Ile	Ala	Leu	Asp	Pro	Arg	Asp	Gly	Ile	Leu	Phe	Trp	Thr	Asp	Trp	Asp		
	1395						1400					1405					
Ala	Ser	Leu	Pro	Arg	Ile	Glu	Ala	Ala	Ser	Met	Ser	Gly	Ala	Gly	Arg		
	1410					1415						1420					

Gly Val Gly Ser Phe Leu Leu Tyr Ser Val His Glu Gly Ile Arg Gly	1890	1895	1900
Ile Pro Leu Asp Pro Asn Asp Lys Ser Asp Ala Leu Val Pro Val Ser	1905	1910	1915
Gly Thr Ser Leu Ala Val Gly Ile Asp Phe His Ala Glu Asn Asp Thr	1925	1930	1935
Ile Tyr Trp Val Asp Met Gly Leu Ser Thr Ile Ser Arg Ala Lys Arg	1940	1945	1950
Asp Gln Thr Trp Arg Glu Asp Val Val Thr Asn Gly Ile Gly Arg Val	1955	1960	1965
Glu Gly Ile Ala Val Asp Trp Ile Ala Gly Asn Ile Tyr Trp Thr Asp	1970	1975	1980
Gln Gly Phe Asp Val Ile Glu Val Ala Arg Leu Asn Gly Ser Phe Arg	1985	1990	1995
Tyr Val Val Ile Ser Gln Gly Leu Asp Lys Pro Arg Ala Ile Thr Val	2005	2010	2015
His Pro Glu Lys Gly Tyr Leu Phe Trp Thr Glu Trp Gly His Tyr Pro	2020	2025	2030
Arg Ile Glu Arg Ser Arg Leu Asp Gly Thr Glu Arg Val Val Leu Val	2035	2040	2045
Asn Val Ser Ile Ser Trp Pro Asn Gly Ile Ser Val Asp Tyr Gln Gly	2050	2055	2060
Gly Lys Leu Tyr Trp Cys Asp Ala Arg Met Asp Lys Ile Glu Arg Ile	2065	2070	2075
Asp Leu Glu Thr Gly Glu Asn Arg Glu Val Val Leu Ser Ser Asn Asn	2085	2090	2095
Met Asp Met Phe Ser Val Ser Val Phe Glu Asp Phe Ile Tyr Trp Ser	2100	2105	2110
Asp Arg Thr His Ala Asn Gly Ser Ile Lys Arg Gly Cys Lys Asp Asn	2115	2120	2125
Ala Thr Asp Ser Val Pro Leu Arg Thr Gly Ile Gly Val Gln Leu Lys	2130	2135	2140
Asp Ile Lys Val Phe Asn Arg Asp Arg Gln Lys Gly Thr Asn Val Cys	2145	2150	2155
Ala Val Ala Asn Gly Gly Cys Gln Gln Leu Cys Leu Tyr Arg Gly Gly	2165	2170	2175
Gly Gln Arg Ala Cys Ala Cys Ala His Gly Met Leu Ala Glu Asp Gly	2180	2185	2190
Ala Ser Cys Arg Glu Tyr Ala Gly Tyr Leu Leu Tyr Ser Glu Arg Thr	2195	2200	2205
Ile Leu Lys Ser Ile His Leu Ser Asp Glu Arg Asn Leu Asn Ala Pro	2210	2215	2220
Val Gln Pro Phe Glu Asp Pro Glu His Met Lys Asn Val Ile Ala Leu	2225	2230	2235
Ala Phe Asp Tyr Arg Ala Gly Thr Ser Pro Gly Thr Pro Asn Arg Ile	2245	2250	2255
Phe Phe Ser Asp Ile His Phe Gly Asn Ile Gln Gln Ile Asn Asp Asp	2260	2265	2270
Gly Ser Gly Arg Thr Thr Ile Val Glu Asn Val Gly Ser Val Glu Gly	2275	2280	2285
Leu Ala Tyr His Arg Gly Trp Asp Thr Leu Tyr Trp Thr Ser Tyr Thr	2290	2295	2300
Thr Ser Thr Ile Thr Arg His Thr Val Asp Gln Thr Arg Pro Gly Ala	2305	2310	2315
Phe Glu Arg Glu Thr Val Ile Thr Met Ser Gly Asp Asp His Pro Arg	2325	2330	2335
Ala Phe Val Leu Asp Glu Cys Gln Asn Leu Met Phe Trp Thr Asn Trp	2340	2345	2350

Asn	Glu	Leu	His	Pro	Ser	Ile	Met	Arg	Ala	Ala	Leu	Ser	Gly	Ala	Asn	2355	2360	2365
Val	Leu	Thr	Leu	Ile	Glu	Lys	Asp	Ile	Arg	Thr	Pro	Asn	Gly	Leu	Ala	2370	2375	2380
Ile	Asp	His	Arg	Ala	Glu	Lys	Leu	Tyr	Phe	Ser	Asp	Ala	Thr	Leu	Asp	2385	2390	2395
Lys	Ile	Glu	Arg	Cys	Glu	Tyr	Asp	Gly	Ser	His	Arg	Tyr	Val	Ile	Leu	2405	2410	2415
Lys	Ser	Glu	Pro	Val	His	Pro	Phe	Gly	Leu	Ala	Val	Tyr	Gly	Glu	His	2420	2425	2430
Ile	Phe	Trp	Thr	Asp	Trp	Val	Arg	Arg	Ala	Val	Gln	Arg	Ala	Asn	Lys	2435	2440	2445
Tyr	Val	Gly	Ser	Asp	Met	Lys	Leu	Leu	Arg	Val	Asp	Ile	Pro	Gln	Gln	2450	2455	2460
Pro	Met	Gly	Ile	Ile	Ala	Val	Ala	Asn	Asp	Thr	Asn	Ser	Cys	Glu	Leu	2465	2470	2475
Ser	Pro	Cys	Arg	Ile	Asn	Asn	Gly	Gly	Cys	Gln	Asp	Leu	Cys	Leu	Leu	2485	2490	2495
Thr	His	Gln	Gly	His	Val	Asn	Cys	Ser	Cys	Arg	Gly	Gly	Arg	Ile	Leu	2500	2505	2510
Gln	Glu	Asp	Phe	Thr	Cys	Arg	Ala	Val	Asn	Ser	Ser	Cys	Arg	Ala	Gln	2515	2520	2525
Asp	Glu	Phe	Glu	Cys	Ala	Asn	Gly	Glu	Cys	Ile	Ser	Phe	Ser	Leu	Thr	2530	2535	2540
Cys	Asp	Gly	Val	Ser	His	Cys	Lys	Asp	Lys	Ser	Asp	Glu	Lys	Pro	Ser	2545	2550	2555
Tyr	Cys	Asn	Ser	Arg	Arg	Cys	Lys	Lys	Thr	Phe	Arg	Gln	Cys	Asn	Asn	2565	2570	2575
Gly	Arg	Cys	Val	Ser	Asn	Met	Leu	Trp	Cys	Asn	Gly	Val	Asp	Tyr	Cys	2580	2585	2590
Gly	Asp	Gly	Ser	Asp	Glu	Ile	Pro	Cys	Asn	Lys	Thr	Ala	Cys	Gly	Val	2595	2600	2605
Gly	Glu	Phe	Arg	Cys	Arg	Asp	Gly	Ser	Cys	Ile	Gly	Asn	Ser	Ser	Arg	2610	2615	2620
Cys	Asn	Gln	Phe	Val	Asp	Cys	Glu	Asp	Ala	Ser	Asp	Glu	Met	Asn	Cys	2625	2630	2635
Ser	Ala	Thr	Asp	Cys	Ser	Ser	Tyr	Phe	Arg	Leu	Gly	Val	Lys	Gly	Val	2645	2650	2655
Leu	Phe	Gln	Pro	Cys	Glu	Arg	Thr	Ser	Leu	Cys	Tyr	Ala	Pro	Ser	Trp	2660	2665	2670
Val	Cys	Asp	Gly	Ala	Asn	Asp	Cys	Gly	Asp	Tyr	Ser	Asp	Glu	Arg	Asp	2675	2680	2685
Cys	Pro	Gly	Val	Lys	Arg	Pro	Arg	Cys	Pro	Leu	Asn	Tyr	Phe	Ala	Cys	2690	2695	2700
Pro	Ser	Gly	Arg	Cys	Ile	Pro	Met	Ser	Trp	Thr	Cys	Asp	Lys	Glu	Asp	2705	2710	2715
Asp	Cys	Glu	Asn	Gly	Glu	Asp	Glu	Thr	His	Cys	Asn	Lys	Phe	Cys	Ser	2725	2730	2735
Glu	Ala	Gln	Phe	Glu	Cys	Gln	Asn	His	Arg	Cys	Ile	Ser	Lys	Gln	Trp	2740	2745	2750
Leu	Cys	Asp	Gly	Ser	Asp	Asp	Cys	Gly	Asp	Gly	Ser	Asp	Glu	Ala	Ala	2755	2760	2765
His	Cys	Glu	Gly	Lys	Thr	Cys	Gly	Pro	Ser	Ser	Phe	Ser	Cys	Pro	Gly	2770	2775	2780
Thr	His	Val	Cys	Val	Pro	Glu	Arg	Trp	Leu	Cys	Asp	Gly	Asp	Lys	Asp	2785	2790	2795
Cys	Thr	Asp	Gly	Ala	Asp	Glu	Ser	Val	Thr	Ala	Gly	Cys	Leu	Tyr	Asn	2805	2810	2815

Ser Thr Cys Asp Asp Arg Glu Phe Met Cys Gln Asn Arg Leu Cys Ile
 2820 2825 2830
 Pro Lys His Phe Val Cys Asp His Asp Arg Asp Cys Ala Asp Gly Ser
 2835 2840 2845
 Asp Glu Ser Pro Glu Cys Glu Tyr Pro Thr Cys Gly Pro Asn Glu Phe
 2850 2855 2860
 Arg Cys Ala Asn Gly Arg Cys Leu Ser Ser Arg Gln Trp Glu Cys Asp
 2865 2870 2875 2880
 Gly Glu Asn Asp Cys His Asp His Ser Asp Glu Ala Pro Lys Asn Pro
 2885 2890 2895
 His Cys Thr Ser Pro Glu His Lys Cys Asn Ala Ser Ser Gln Phe Leu
 2900 2905 2910
 Cys Ser Ser Gly Arg Cys Val Ala Glu Ala Leu Leu Cys Asn Gly Gln
 2915 2920 2925
 Asp Asp Cys Gly Asp Gly Ser Asp Glu Arg Gly Cys His Val Asn Glu
 2930 2935 2940
 Cys Leu Ser Arg Lys Leu Ser Gly Cys Ser Gln Asp Cys Glu Asp Leu
 2945 2950 2955 2960
 Lys Ile Gly Phe Lys Cys Arg Cys Arg Pro Gly Phe Arg Leu Lys Asp
 2965 2970 2975
 Asp Gly Arg Thr Cys Ala Asp Leu Asp Glu Cys Ser Thr Thr Phe Pro
 2980 2985 2990
 Cys Ser Gln Leu Cys Ile Asn Thr His Gly Ser Tyr Lys Cys Leu Cys
 2995 3000 3005
 Val Glu Gly Tyr Ala Pro Arg Gly Gly Asp Pro His Ser Cys Lys Ala
 3010 3015 3020
 Val Thr Asp Glu Glu Pro Phe Leu Ile Phe Ala Asn Arg Tyr Tyr Leu
 3025 3030 3035 3040
 Arg Lys Leu Asn Leu Asp Gly Ser Asn Tyr Thr Leu Leu Lys Gln Gly
 3045 3050 3055
 Leu Asn Asn Ala Val Ala Leu Ala Phe Asp Tyr Arg Glu Gln Met Ile
 3060 3065 3070
 Tyr Trp Thr Gly Val Thr Thr Gln Gly Ser Met Ile Arg Arg Met His
 3075 3080 3085
 Leu Asn Gly Ser Asn Val Gln Val Leu His Arg Thr Gly Leu Ser Asn
 3090 3095 3100
 Pro Asp Gly Leu Ala Val Asp Trp Val Gly Gly Asn Leu Tyr Trp Cys
 3105 3110 3115 3120
 Asp Lys Gly Arg Asp Thr Ile Glu Val Ser Lys Leu Asn Gly Ala Tyr
 3125 3130 3135
 Arg Thr Val Leu Val Ser Ser Gly Leu Arg Glu Pro Arg Ala Leu Val
 3140 3145 3150
 Val Asp Val Gln Asn Gly Tyr Leu Tyr Trp Thr Asp Trp Gly Asp His
 3155 3160 3165
 Ser Leu Ile Gly Arg Ile Gly Met Asp Gly Ser Gly Arg Ser Ile Ile
 3170 3175 3180
 Val Asp Thr Lys Ile Thr Trp Pro Asn Gly Leu Thr Val Asp Tyr Val
 3185 3190 3195 3200
 Thr Glu Arg Ile Tyr Trp Ala Asp Ala Arg Glu Asp Tyr Ile Glu Phe
 3205 3210 3215
 Ala Ser Leu Asp Gly Ser Asn Arg His Val Val Leu Ser Gln Asp Ile
 3220 3225 3230
 Pro His Ile Phe Ala Leu Thr Leu Phe Glu Asp Tyr Val Tyr Trp Thr
 3235 3240 3245
 Asp Trp Glu Thr Lys Ser Ile Asn Arg Ala His Lys Thr Thr Gly Ala
 3250 3255 3260
 Asn Lys Thr Leu Leu Ile Ser Thr Leu His Arg Pro Met Asp Leu His
 3265 3270 3275 3280

Val	Phe	His	Ala	Leu	Arg	Gln	Pro	Asp	Val	Pro	Asn	His	Pro	Cys	Lys	
				3285					3290					3295		
Val	Asn	Asn	Gly	Gly	Cys	Ser	Asn	Leu	Cys	Leu	Leu	Ser	Pro	Gly	Gly	
			3300					3305					3310			
Gly	His	Lys	Cys	Ala	Cys	Pro	Thr	Asn	Phe	Tyr	Leu	Gly	Gly	Asp	Gly	
		3315					3320					3325				
Arg	Thr	Cys	Val	Ser	Asn	Cys	Thr	Ala	Ser	Gln	Phe	Val	Cys	Lys	Asn	
	3330					3335					3340					
Asp	Lys	Cys	Ile	Pro	Phe	Trp	Trp	Lys	Cys	Asp	Thr	Glu	Asp	Asp	Cys	
3345					3350					3355					3360	
Gly	Asp	His	Ser	Asp	Glu	Pro	Pro	Asp	Cys	Pro	Glu	Phe	Lys	Cys	Arg	
			3365					3370					3375			
Pro	Gly	Gln	Phe	Gln	Cys	Ser	Thr	Gly	Ile	Cys	Thr	Asn	Pro	Ala	Phe	
			3380					3385					3390			
Ile	Cys	Asp	Gly	Asp	Asn	Asp	Cys	Gln	Asp	Asn	Ser	Asp	Glu	Ala	Asn	
	3395					3400					3405					
Cys	Asp	Ile	His	Val	Cys	Leu	Pro	Ser	Gln	Phe	Lys	Cys	Thr	Asn	Thr	
	3410					3415					3420					
Asn	Arg	Cys	Ile	Pro	Gly	Ile	Phe	Arg	Cys	Asn	Gly	Gln	Asp	Asn	Cys	
3425				3430				3435							3440	
Gly	Asp	Gly	Glu	Asp	Glu	Arg	Asp	Cys	Pro	Glu	Val	Thr	Cys	Ala	Pro	
			3445					3450					3455			
Asn	Gln	Phe	Gln	Cys	Ser	Ile	Thr	Lys	Arg	Cys	Ile	Pro	Arg	Val	Trp	
		3460					3465					3470				
Val	Cys	Asp	Arg	Asp	Asn	His	Cys	Val	Asp	Gly	Ser	Asp	Glu	Pro	Ala	
	3475					3480					3485					
Asn	Cys	Thr	Gln	Met	Thr	Cys	Gly	Val	Asp	Glu	Phe	Arg	Cys	Lys	Asp	
	3490				3495				3500							
Ser	Gly	Arg	Cys	Ile	Pro	Ala	Arg	Trp	Lys	Cys	Asp	Gly	Glu	Asp	Asp	
3505				3510				3515							3520	
Cys	Gly	Asp	Gly	Ser	Asp	Glu	Pro	Lys	Glu	Glu	Cys	Asp	Glu	Arg	Thr	
			3525					3530					3535			
Cys	Glu	Pro	Tyr	Gln	Phe	Arg	Cys	Lys	Asn	Asn	Arg	Cys	Val	Pro	Gly	
		3540						3545					3550			
Arg	Trp	Gln	Cys	Asp	Tyr	Asp	Asn	Asp	Cys	Gly	Asp	Asn	Ser	Asp	Glu	
	3555					3560						3565				
Glu	Ser	Cys	Thr	Pro	Arg	Pro	Cys	Ser	Glu	Ser	Glu	Phe	Phe	Cys	Ala	
	3570					3575					3580					
Asn	Gly	Arg	Cys	Ile	Ala	Gly	Arg	Trp	Lys	Cys	Asp	Gly	Asp	His	Asp	
3585				3590				3595							3600	
Cys	Ala	Asp	Gly	Ser	Asp	Glu	Lys	Asp	Cys	Thr	Pro	Arg	Cys	Asp	Met	
			3605					3610					3615			
Asp	Gln	Phe	Gln	Cys	Lys	Ser	Gly	His	Cys	Ile	Pro	Leu	Arg	Trp	Pro	
		3620					3625					3630				
Cys	Asp	Ala	Asp	Ala	Asp	Cys	Met	Asp	Gly	Ser	Asp	Glu	Glu	Ala	Cys	
	3635					3640					3645					
Gly	Thr	Gly	Val	Arg	Thr	Cys	Pro	Leu	Asp	Glu	Phe	Gln	Cys	Asn	Asn	
	3650					3655					3660					
Thr	Leu	Cys	Lys	Pro	Leu	Ala	Trp	Lys	Cys	Asp	Gly	Glu	Asp	Asp	Cys	
3665				3670				3675							3680	
Gly	Asp	Asn	Ser	Asp	Glu	Asn	Pro	Glu	Glu	Cys	Ala	Arg	Phe	Ile	Cys	
			3685					3690					3695			
Pro	Pro	Asn	Arg	Pro	Phe	Arg	Cys	Lys	Asn	Asp	Arg	Val	Cys	Leu	Trp	
		3700					3705					3710				
Ile	Gly	Arg	Gln	Cys	Asp	Gly	Val	Asp	Asn	Cys	Gly	Asp	Gly	Thr	Asp	
	3715					3720					3725					
Glu	Glu	Asp	Cys	Glu	Pro	Pro	Thr	Ala	Gln	Asn	Pro	His	Cys	Lys	Asp	
	3730					3735					3740					

Lys Lys Glu Phe Leu Cys Arg Asn Gln Arg Cys Leu Ser Ser Ser Leu
 3745 3750 3755 3760
 Arg Cys Asn Met Phe Asp Asp Cys Gly Asp Gly Ser Asp Glu Glu Asp
 3765 3770 3775
 Cys Ser Ile Asp Pro Lys Leu Thr Ser Cys Ala Thr Asn Ala Ser Met
 3780 3785 3790
 Cys Gly Asp Glu Ala Arg Cys Val Arg Thr Glu Lys Ala Ala Tyr Cys
 3795 3800 3805
 Ala Cys Arg Ser Gly Phe His Thr Val Pro Gly Gln Pro Gly Cys Gln
 3810 3815 3820
 Asp Ile Asn Glu Cys Leu Arg Phe Gly Thr Cys Ser Gln Leu Trp Asn
 3825 3830 3835 3840
 Lys Pro Lys Gly Gly His Leu Cys Ser Cys Ala Arg Asn Phe Met Lys
 3845 3850 3855
 Thr His Asn Thr Cys Lys Ala Glu Gly Ser Glu Tyr Gln Val Leu Tyr
 3860 3865 3870
 Ile Ala Asp Asp Asn Glu Ile Arg Ser Leu Phe Pro Gly His Pro His
 3875 3880 3885
 Ser Ala Tyr Glu Gln Thr Phe Gln Gly Asp Glu Ser Val Arg Ile Asp
 3890 3895 3900
 Ala Met Asp Val His Val Lys Ala Gly Arg Val Tyr Trp Thr Asn Trp
 3905 3910 3915 3920
 His Thr Gly Thr Ile Ser Tyr Arg Ser Leu Pro Pro Ala Ala Pro Pro
 3925 3930 3935
 Thr Thr Ser Asn Arg His Arg Arg Gln Ile Asp Arg Gly Val Thr His
 3940 3945 3950
 Leu Asn Ile Ser Gly Leu Lys Met Pro Arg Gly Ile Ala Ile Asp Trp
 3955 3960 3965
 Val Ala Gly Asn Val Tyr Trp Thr Asp Ser Gly Arg Asp Val Ile Glu
 3970 3975 3980
 Val Ala Gln Met Lys Gly Glu Asn Arg Lys Thr Leu Ile Ser Gly Met
 3985 3990 3995 4000
 Ile Asp Glu Pro His Ala Ile Val Val Asp Pro Leu Arg Gly Thr Met
 4005 4010 4015
 Tyr Trp Ser Asp Trp Gly Asn His Pro Lys Ile Glu Thr Ala Ala Met
 4020 4025 4030
 Asp Gly Thr Leu Arg Glu Thr Leu Val Gln Asp Asn Ile Gln Trp Pro
 4035 4040 4045
 Thr Gly Leu Ala Val Asp Tyr His Asn Glu Arg Leu Tyr Trp Ala Asp
 4050 4055 4060
 Ala Lys Leu Ser Val Ile Gly Ser Ile Arg Leu Asn Gly Thr Asp Pro
 4065 4070 4075 4080
 Ile Val Ala Ala Asp Ser Lys Arg Gly Leu Ser His Pro Phe Ser Ile
 4085 4090 4095
 Asp Val Phe Glu Asp Tyr Ile Tyr Gly Val Thr Tyr Ile Asn Asn Arg
 4100 4105 4110
 Val Phe Lys Ile His Lys Phe Gly His Ser Pro Leu Tyr Asn Leu Thr
 4115 4120 4125
 Gly Gly Leu Ser His Ala Ser Asp Val Val Leu Tyr His Gln His Lys
 4130 4135 4140
 Gln Pro Glu Val Thr Asn Pro Cys Asp Arg Lys Lys Cys Glu Trp Leu
 4145 4150 4155 4160
 Cys Leu Leu Ser Pro Ser Gly Pro Val Cys Thr Cys Pro Asn Gly Lys
 4165 4170 4175
 Arg Leu Asp Asn Gly Thr Cys Val Pro Val Pro Ser Pro Thr Pro Pro
 4180 4185 4190
 Pro Asp Ala Pro Arg Pro Gly Thr Cys Thr Leu Gln Cys Phe Asn Gly
 4195 4200 4205

Gly Ser Cys Phe Leu Asn Ala Arg Arg Gln Pro Lys Cys Arg Cys Gln
 4210 4215 4220
 Pro Arg Tyr Thr Gly Asp Lys Cys Glu Leu Asp Gln Cys Trp Glu Tyr
 4225 4230 4235 4240
 Cys His Asn Gly Gly Thr Cys Ala Ala Ser Pro Ser Gly Met Pro Thr
 4245 4250 4255
 Cys Arg Cys Pro Thr Gly Phe Thr Gly Pro Lys Cys Thr Ala Gln Val
 4260 4265 4270
 Cys Ala Gly Tyr Cys Ser Asn Asn Ser Thr Cys Thr Val Asn Gln Gly
 4275 4280 4285
 Asn Gln Pro Gln Cys Arg Cys Leu Pro Gly Phe Leu Gly Asp Arg Cys
 4290 4295 4300
 Gln Tyr Arg Gln Cys Ser Gly Phe Cys Glu Asn Phe Gly Thr Cys Gln
 4305 4310 4315 4320
 Met Ala Ala Asp Gly Ser Arg Gln Cys Arg Cys Thr Val Tyr Phe Glu
 4325 4330 4335
 Gly Pro Arg Cys Glu Val Asn Lys Cys Ser Arg Cys Leu Gln Gly Ala
 4340 4345 4350
 Cys Val Val Asn Lys Gln Thr Gly Asp Val Thr Cys Asn Cys Thr Asp
 4355 4360 4365
 Gly Arg Val Ala Pro Ser Cys Leu Thr Cys Ile Asp His Cys Ser Asn
 4370 4375 4380
 Gly Gly Ser Cys Thr Met Asn Ser Lys Met Met Pro Glu Cys Gln Cys
 4385 4390 4395 4400
 Pro Pro His Met Thr Gly Pro Arg Cys Gln Glu Gln Val Val Ser Gln
 4405 4410 4415
 Gln Gln Pro Gly His Met Ala Ser Ile Leu Ile Pro Leu Leu Leu Leu
 4420 4425 4430
 Leu Leu Leu Leu Leu Val Ala Gly Val Val Phe Trp Tyr Lys Arg Arg
 4435 4440 4445
 Val Arg Gly Ala Lys Gly Phe Gln His Gln Arg Met Thr Asn Gly Ala
 4450 4455 4460
 Met Asn Val Glu Ile Gly Asn Pro Thr Tyr Lys Met Tyr Glu Gly Gly
 4465 4470 4475 4480
 Glu Pro Asp Asp Val Gly Gly Leu Leu Asp Ala Asp Phe Ala Leu Asp
 4485 4490 4495
 Pro Asp Lys Pro Thr Asn Phe Thr Asn Pro Val Tyr Ala Thr Leu Tyr
 4500 4505 4510
 Met Gly Gly His Gly Ser Arg His Ser Leu Ala Ser Thr Asp Glu Lys
 4515 4520 4525
 Arg Glu Leu Leu Gly Arg Gly Pro Glu Asp Glu Ile Gly Asp Pro Leu
 4530 4535 4540
 Ala
 4545

<210> 3
 <211> 4577
 <212> DNA
 <213> Homo sapiens

<400> 3
 gctacaatcc atctggtctc ctccagctcc ttcttttctgc aacatgggga agaacaaact 60
 ccttcatcca agtctggttc ttctcctctt ggctcctctg cccacagacg cctcagtcctc 120
 tggaaaaccg cagtatatgg ttctgggtccc ctccctgctc cacactgaga ccactgagaa 180
 gggctgtgtc cttctgagct acctgaatga gacagtgtact gtaagtgtct ccttggagtc 240
 tgtcagggga aacaggagcc tcttcactga cctggaggcg gagaatgacg tactccactg 300
 tgtcgcttcc gctgtcccaa agtcttcatc caatgaggag gtaatgttcc tcaactgtcca 360
 agtgaaagga ccaacccaag aatttaagaa gcggaccaca gtgatgggta agaacgagga 420

cagtctggtc	tttgtccaga	cagacaaatc	aatctacaaa	ccagggcaga	cagtgaaatt	480
tcgtgttggtc	tccatggatg	aaaactttca	ccccctgaat	gagttgattc	cactagtata	540
cattcaggat	cccaaaggaa	atcgcacgc	acaatggcag	agtttccagt	tagaggggtg	600
cctcaagcaa	ttttcttttc	ccctctcatc	agagcccttc	cagggctcct	acaaggtggt	660
ggtacagaag	aaatcaggtg	gaaggacaga	gcaccctttc	accgtggagg	aatttggttct	720
tcccaagttt	gaagtacaag	taacagtgcc	aaagataatc	accatcttgg	aagaagagat	780
gaatgtatca	gtgtgtggcc	tatacacata	tgggaagcct	gtccctggac	atgtgactgt	840
gagcatttgc	agaaagtata	gtgacgcttc	cgactgccac	ggtgaagatt	cacaggcttt	900
ctgtgagaaa	ttcagtggac	agctaaacag	ccatggctgc	ttctatcagc	aagtaaaaac	960
caaggtcttc	cagctgaaga	ggaaggagta	tgaaatgaaa	cttcacactg	aggcccagat	1020
ccaagaagaa	ggaacagtgg	tggaattgac	tggaaggcag	tccagtgaaa	tcacaagaac	1080
cataaccaaa	ctctcatttg	tgaaagtgga	ctcacacttt	cgacagggaa	ttcccttctt	1140
tgggcaggtg	cgcttagtag	atgggaaagg	cgctccctata	ccaaataaag	tcatattcat	1200
cagaggaaat	gaagcaaact	attactccaa	tgctaccacg	gatgagcatg	gccttgtaca	1260
gttctctatc	aacaccacca	acgttatggg	tacctctctt	actgttaggg	tcaattacaa	1320
ggatcgtagt	ccctgttacg	gctaccagtg	ggtgtcagaa	gaacacgaag	aggcacatca	1380
cactgcttat	cttgtgttct	ccccaagcaa	gagctttgtc	caccttgagc	ccatgtctca	1440
tgaactacc	tgtggccata	ctcagacagt	ccaggcacat	tatattctga	atggaggcac	1500
cctgctgggg	ctgaagaagc	tctcctttta	ttatctgata	atggcaaagg	gaggcattgt	1560
ccgaactggg	actcatggac	tgcttgtgaa	gcaggaagac	atgaagggcc	atttttccat	1620
ctcaatccct	gtgaagtcag	acattgctcc	tgctcgctcg	ttgctcatct	atgctgtttt	1680
acctaccggg	gacgtgattg	gggattctgc	aaaatatgat	gttgaaaatt	gtctggccaa	1740
caaggtggat	ttgagcttca	gcccatcaca	aagtctccca	gcctcacacg	cccacctgcg	1800
agtcacagcg	gctcctcagt	ccgtctgcgc	cctccgtgct	gtggaccaa	gcgtgctgct	1860
catgaagcct	gatgctgagc	tctcggcgtc	ctcggtttac	aacctgctac	cagaaaagga	1920
cctcactggc	ttccctgggc	ctttgaatga	ccaggacgat	gaagactgca	tcaatcgtca	1980
taatgtctat	attaatggaa	tcacatatat	tccagtatca	agtacaaatg	aaaaggatat	2040
gtacagcttc	ctagaggaca	tgggcttaaa	ggcattcacc	aactcaaaga	ttcgtaaacc	2100
caaaatgtgt	ccacagcttc	aacagtatga	aatgcatgga	cctgaaggtc	tacgtgtagg	2160
tttttatgag	tcagatgtaa	tgggaagagg	ccatgcacgc	ctggtgcatg	ttgaagagcc	2220
tcacacggag	accgtacgaa	agtacttccc	tgagacatgg	atctgggatt	tgggtggtggt	2280
aaactcagca	ggggtggctg	aggtaggagt	aacagtcctt	gacaccatca	ccgagtggaa	2340
ggcagggggc	ttctgcctgt	ctgaagatgc	tggacttggg	atctcttcca	ctgcctctct	2400
ccgagccttc	cagcccttct	ttgtggagct	tacaatgcct	tactctgtga	ttcgtggaga	2460
ggccttcaca	ctcaaggcca	cggtcctaaa	ctaccttccc	aatgcatcc	gggtcagtgt	2520
gcagctggaa	gcctctcccc	ccttccttgc	tgtcccagtg	gagaaggaa	aagcgctca	2580
ctgcatctgt	gcaaacgggc	ggcaaactgt	gtcctgggca	gtaaccccaa	agtcattagg	2640
aaatgtgaat	ttcactgtga	gcgcagaggc	actagagtct	caagagctgt	gtgggactga	2700
ggtgccttca	gttctgaac	acggaaggaa	agacacagtc	atcaagcctc	tgttggttga	2760
acctgaagga	ctagagaagg	aaacaacatt	caactcccta	ctttgtccat	caggtggtga	2820
ggtttctgaa	gaattatccc	tgaaactgcc	accaaattgtg	gtagaagaat	ctgcccagagc	2880
ttctgtctca	gttttgggag	acatattagg	ctctgccatg	caaaacacac	aaaatcttct	2940
ccagatgccc	tatggctgtg	gagagcagaa	tatggctctc	tttgcctcta	acatctatgt	3000
actggattat	ctaaatgaaa	cacagcagct	tactccagag	gtcaagtcca	aggccattgg	3060
ctatctcaac	actggttacc	agagacagtt	gaactacaaa	cactatgatg	gctcctacag	3120
cacctttggg	gagcgatatg	gcaggaacca	gggcaacacc	tggctcacag	cctttgttct	3180
gaagactttt	gccaagctc	gagcctacat	cttcacgat	gaagcacaca	ttacccaagc	3240
cctcatatgg	ctctcccaga	ggcagaagga	caatggctgt	ttcaggagct	ctgggtcact	3300
gctcaacaat	gccataaagg	gaggagtaga	agatgaagtg	accctctccg	cctatatcac	3360
catcgccctt	ctggagattc	ctctcacagt	cactcacctt	gttgctccga	atgccctggt	3420
ttgcctggag	tcagcctgga	agacagcaca	agaagggggac	catggcagcc	atgtatatac	3480
caaagcactg	ctggcctatg	cttttgccct	ggcaggtaac	caggacaaga	ggaagggaagt	3540
actcaagtca	cttaatgagg	aagctgtgaa	gaaagacaa	tctgtccatt	gggagcgccc	3600
tcagaaaccc	aaggcaccag	tggggcattt	ttacgaaccc	caggctccct	ctgctgaggt	3660
ggagatgaca	tcctatgtgc	tcctcgctta	tctcacggcc	cagccagccc	caacctcgga	3720
ggacctgacc	tctgcaacca	acatcgtgaa	gtggatcacg	aagcagcaga	atgcccaggg	3780
cggtttctcc	tccaccaggg	acacagtggg	ggctctccat	gctctgtcca	aatatggagc	3840
cgccacattt	accaqgactg	ggaaggctgc	ac			

ttccagcaaa	ttccaagtgg	acaacaacaa	tcgcctgtta	ctgcagcagg	tctcattgcc	3960
agagctgcct	ggggaataca	gcatgaaagt	gacaggagaa	ggatgtgtct	acctccagac	4020
ctccttgaaa	tacaatatct	tcccagaaaa	ggaagagttc	ccctttgtct	taggagtgc	4080
gactctgcct	caaacttgtg	atgaacccaa	agcccacacc	agcttccaaa	tctccctaag	4140
tgtcagttac	acagggagcc	gctctgcctc	caacatggcg	atcggtgatg	tgaagatggg	4200
ctctggcttc	attcccctga	agccaacagt	gaaaatgctt	gaaagatcta	accatgtgag	4260
ccggacagaa	gtcagcagca	accatgtctt	gatttacctt	gataaggtgt	caaatacagac	4320
actgagcttg	ttcttcacgg	ttctgcaaga	tgtcccagta	agagatctca	aaccagccat	4380
agtgaaagtc	tatgattact	acgagacgga	tgagtttgca	atcgctgagt	acaatgctcc	4440
ttgcagcaaa	gatcttgga	atgcttgaag	accacaaggc	tgaaaagtgc	tttgctggag	4500
tcctgttctc	tgagctccac	agaagacacg	tgtttttgta	tcttttaaaga	cttgatgaat	4560
aaacactttt	tctgggtc					4577

<210> 4
 <211> 4422
 <212> DNA
 <213> Homo sapiens

<400> 4						
atggggaaga	acaaactcct	tcateccaagt	ctgggttcttc	tcctcttggt	cctcctgccc	60
acagacgcct	cagtctctgg	aaaaccgcag	tatatgggtc	tggtcccttc	cctgctccac	120
actgagacca	ctgagaagg	ctgtgtcctt	ctgagctacc	tgaatgagac	agtgactgta	180
agtgtctcct	tggagtctgt	caggggaaac	aggagcctct	tcactgacct	ggaggcggag	240
aatgacgtac	tccactgtgt	cgccttcgct	gtcccaaagt	cttcatccaa	tgaggaggta	300
atgttcctca	ctgtccaagt	gaaaggacca	acccaagaat	ttaagaagcg	gaccacagtg	360
atggttaaga	acgaggacag	tctgggtcttt	gtccagacag	acaaatcaat	ctacaaacca	420
gggcagacag	tgaaatttcg	tgttgctctcc	atggatgaaa	actttcaccc	cctgaatgag	480
ttgattccac	tagtatacat	tcaggatccc	aaaggaaatc	gcatcgcaca	atggcagagt	540
ttccagttag	agggtggcct	caagcaattt	tcttttcccc	tctcatcaga	gcccttccag	600
ggctcctaca	agggtgggtg	acagaagaaa	tcagggtggaa	ggacagagca	ccctttcacc	660
gtggagggaat	ttgttcttcc	caagtttgaa	gtacaagtaa	cagtgccaaa	gataatcacc	720
atcttggaag	aagagatgaa	tgtatcagtg	tgtggcctat	acacatatgg	gaagcctgtc	780
cctggacatg	tgactgtgag	catttgcaga	aagtatagtg	acgcttccga	ctgccacggg	840
gaagattcac	aggctttctg	tgagaaattc	agtggacagc	taaacagcca	tggtctgttc	900
tatcagcaag	taaaaacca	ggtcttccag	ctgaagagga	aggagtatga	aatgaaactt	960
cacactgagg	cccagatcca	agaagaagga	acagtgggtg	aattgactgg	aaggcagtc	1020
agtgaatca	caagaaccat	aaccaaactc	tcattttgtga	aagtggactc	acactttcga	1080
caggggaattc	ccttcttttg	gcagggtgcg	ctagtagatg	ggaaaggcgt	ccctatacca	1140
aataaagtca	tattcatcag	aggaaatgaa	gcaaactatt	actccaatgc	taccacggat	1200
gagcatggcc	ttgtacagtt	ctctatcaac	accaccaacg	ttatgggtac	ctctcttact	1260
gttagggtea	attacaagga	tcgtagtccc	tgttacggct	accagtgggt	gtcagaagaa	1320
cacgaagagg	cacatcacac	tgcttatctt	gtgttctccc	caagcaagag	ctttgtccac	1380
cttgagccca	tgtctcatga	actaccctgt	ggccatactc	agacagtcca	ggcacattat	1440
attctgaatg	gaggcaccct	gctggggctg	aagaagctct	cctttttatta	tctgataatg	1500
gcaaaggagg	gcattgtccg	aactgggact	catggactgc	ttgtgaagca	ggaagacatg	1560
aagggccatt	tttccatctc	aatccctgtg	aagtcagaca	ttgtcctgtg	cgctcggttg	1620
ctcatctatg	ctgtttttacc	taccggggac	gtgattgggg	attctgcaaa	atatgatgtt	1680
gaaaattgtc	tggccaacaa	ggtggatttg	agcttcagcc	catcacaag	tctcccagcc	1740
tcacacgccc	acctgcgagt	cacagcggct	cctcagtcct	tctgcgcctc	ccgtgctgtg	1800
gaccaaagcg	tgctgctcat	gaagcctgat	gctgagctct	cggcgtcctc	ggtttacaac	1860
ctgctaccag	aaaaggacct	cactggcttc	cctgggcctt	tgaatgacca	ggacgatgaa	1920
gactgcatca	atcgtcataa	tgtctatatt	aatggaatca	catatactcc	agtatcaagt	1980
acaaatgaaa	aggatatgta	cagcttccta	gaggacatgg	gcttaaaggc	attcaccaac	2040
tcaaagattc	gtaaacccaa	aatgtgtcca	cagcttcaac	agtatgaaat	gcatggacct	2100
gaaggctctac	gtgtagggtt	ttatgagtea	gatgtaatgg	gaagaggcca	tgcacgcctg	2160
gtgcatgttg	aagagcctca	cacggagacc	gtacgaaagt	acttccctga	gacatggatc	2220
tgggatttgg	tggtggtaaa	ctcagcaggg	gtggctgagg	taggagtaac	agtccttgac	2280
accatcaccg	agtggaaggc	aggggccttc	tgccctgtctg	aagatgctgg	acttggtatc	2340

tcttccactg	cctctctccg	agccttccag	cccttctttg	tggagcttac	aatgccttac	2400
tctgtgattc	gtggagaggc	cttcacactc	aaggccacgg	tcctaaacta	ccttcccaaa	2460
tgcacccggg	tcagtgtgca	gctggaagcc	tctcccgcct	tccttgctgt	cccagtgag	2520
aaggaacaag	cgcctcactg	catctgtgca	aacgggcggc	aaactgtgtc	ctgggcagta	2580
accccaaagt	cattaggaaa	tgtgaatttc	actgtgagcg	cagaggcact	agagtctcaa	2640
gagctgtgtg	ggactgaggt	gccttcagtt	cctgaacacg	gaaggaaaga	cacagtcatc	2700
aagcctctgt	tggttgaacc	tgaaggacta	gagaaggaaa	caacattcaa	ctccctactt	2760
tgtccatcag	gtggtgaggt	ttctgaagaa	ttatccctga	aactgccacc	aaatgtggta	2820
gaagaatctg	cccagacttc	tgtctcagtt	ttgggagaca	tattaggctc	tgccatgcaa	2880
aacacacaaa	atcttctcca	gatgccctat	ggctgtggag	agcagaatat	ggtcctcttt	2940
gctcctaaca	tctatgtact	ggattatcta	aatgaaacac	agcagcttac	tccagaggtc	3000
aagtccaagg	ccattggcta	tctcaacact	ggttaccaga	gacagttgaa	ctacaaacac	3060
tatgatggct	cctacagcac	ctttggggag	cgatatggca	ggaaccaggg	caacacctgg	3120
ctcacagcct	ttgttctgaa	gacttttgcc	caagctcgag	cctacatctt	catcgatgaa	3180
gcacacatta	cccaagccct	catatggctc	tcccagaggc	agaaggacaa	tggctgtttc	3240
aggagctctg	ggtcactgct	caacaatgcc	ataaagggag	gagtagaaga	tgaagtgacc	3300
ctctccgcct	atatcaccat	cgcccttctg	gagattcctc	tcacagtcac	tcaccctggt	3360
gtccgcaatg	ccctgttttg	cctggagtca	gcctggaaga	cagcacaaga	aggggaccat	3420
ggcagccatg	tatataccaa	agcactgctg	gcctatgctt	ttgccctggc	aggtaaccag	3480
gacaagagga	aggaagtact	caagtcactt	aatgaggaag	ctgtgaagaa	agacaactct	3540
gtccattggg	agcgccctca	gaaacccaag	gcaccagtgg	ggcattttta	cgaaccccag	3600
gctccctctg	ctgaggtgga	gatgacatcc	tatgtgctcc	tcgcttatct	cacggcccag	3660
ccagccccaa	cctcggagga	cctgacctct	gcaaccaaca	tcgtgaagtg	gatcacgaag	3720
cagcagaatg	cccagggcgg	tttctcctcc	accagggaca	cagtgggtggc	tctccatgct	3780
ctgtccaaat	atggagccgc	cacattttacc	aggactggga	aggctgcaca	ggtgactatc	3840
cagtcttcag	ggacattttc	cagcaaattc	caagtggaca	acaacaatcg	cctgttactg	3900
cagcaggtct	cattgccaga	gctgcctggg	gaatacagca	tgaaagtgac	aggagaagga	3960
tgtgtctacc	tccagacctc	cttgaaatac	aatattctcc	cagaaaagga	agagttcccc	4020
tttgctttag	gagtgcagac	tctgcctcaa	acttgtgatg	aacccaaagc	ccacaccagc	4080
ttccaaatct	ccctaagtgt	cagttacaca	gggagccgct	ctgcctccaa	catggcgatc	4140
gttgatgtga	agatgggtctc	tggcttcatt	cccctgaagc	caacagtgaa	aatgcttgaa	4200
agatctaacc	atgtgagccg	gacagaagtc	agcagcaacc	atgtcttgat	ttaccttgat	4260
aaggtgtcaa	atcagacact	gagcttgttc	ttcacggttc	tgcaagatgt	cccagtaaga	4320
gatctcaaac	cagccatagt	gaaagtctat	gattactacg	agacggatga	gtttgcaatc	4380
gctgaqtaca	atgctccttg	caqcaaaqat	cttgqaaatg	ct		4422

```
<210> 5
<211> 1474
<212> PRT
<213> Homo sapiens
```

<211> 1474

<212> PRT

<213> Homo sapiens

<400> 5

Met 1	Gly	Lys	Asn	Lys 5	Leu	Leu	His	Pro	Ser 10	Leu	Val	Leu	Leu	Leu 15	Leu
Val	Leu	Leu	Pro 20	Thr	Asp	Ala	Ser	Val 25	Ser	Gly	Lys	Pro	Gln 30	Tyr	Met
Val	Leu	Val 35	Pro	Ser	Leu	Leu	His 40	Thr	Glu	Thr	Thr	Glu 45	Lys	Gly	Cys
Val	Leu 50	Leu	Ser	Tyr	Leu	Asn 55	Glu	Thr	Val	Thr	Val 60	Ser	Ala	Ser	Leu
Glu 65	Ser	Val	Arg	Gly	Asn 70	Arg	Ser	Leu	Phe	Thr 75	Asp	Leu	Glu	Ala 80	Glu
Asn	Asp	Val	Leu	His 85	Cys	Val	Ala	Phe 90	Ala	Val	Pro	Lys	Ser 95	Ser	Ser
Asn	Glu	Glu	Val 100	Met	Phe	Leu	Thr	Val 105	Gln	Val	Lys	Gly 110	Pro	Thr	Gln
Glu	Phe	Lys 115	Lys	Arg	Thr	Thr	Val 120	Met	Val	Lys	Asn 125	Glu	Asp	Ser	Leu

1	5	10	15
---	---	----	----

Val Leu Leu Pro Thr Asp Ala Ser Val Ser Gly Lys Pro Gln Tyr Met

20 25 30

Val Leu Val Pro Ser Leu Leu His Thr Glu Thr Thr Glu Lys Gly Cys

Val	Leu	Val	Pro	Ser	Leu	Leu	His	Thr	Glu	Thr	Thr	Glu	Lys	Gly	Cys
		35					40					45			

[illegible]

Val Leu Leu Ser Tyr Leu Asn Glu Thr Val Thr Val Ser Ala Ser Leu
 50 55 60

Gly. Jan. Val. Aug. Gly. Apr. Aug. Jan. Jan. Dec. Thurs. Apr. Jan. Gly. Apr. Gly.

Glu Ser Val Arg Gly Asn Arg Ser Leu Phe Thr Asp Leu Glu Ala Glu
65 70 75 80

65				70				75				80			
Low	Med	Hl	I	Hl	G	Hl	B	B	Hl	P	I	G	G	G	G

Asn Asp Val Leu His Cys Val Ala Phe Ala Val Pro Lys Ser Ser Ser
 65 11 65

85		90		95	
1	1	1	1	1	1
2	2	2	2	2	2
3	3	3	3	3	3
4	4	4	4	4	4
5	5	5	5	5	5
6	6	6	6	6	6
7	7	7	7	7	7
8	8	8	8	8	8
9	9	9	9	9	9
10	10	10	10	10	10
11	11	11	11	11	11
12	12	12	12	12	12
13	13	13	13	13	13
14	14	14	14	14	14
15	15	15	15	15	15
16	16	16	16	16	16
17	17	17	17	17	17
18	18	18	18	18	18
19	19	19	19	19	19
20	20	20	20	20	20
21	21	21	21	21	21
22	22	22	22	22	22
23	23	23	23	23	23
24	24	24	24	24	24
25	25	25	25	25	25
26	26	26	26	26	26
27	27	27	27	27	27
28	28	28	28	28	28
29	29	29	29	29	29
30	30	30	30	30	30
31	31	31	31	31	31
32	32	32	32	32	32
33	33	33	33	33	33
34	34	34	34	34	34
35	35	35	35	35	35
36	36	36	36	36	36
37	37	37	37	37	37
38	38	38	38	38	38
39	39	39	39	39	39
40	40	40	40	40	40
41	41	41	41	41	41
42	42	42	42	42	42
43	43	43	43	43	43
44	44	44	44	44	44
45	45	45	45	45	45
46	46	46	46	46	46
47	47	47	47	47	47
48	48	48	48	48	48
49	49	49	49	49	49
50	50	50	50	50	50
51	51	51	51	51	51
52	52	52	52	52	52
53	53	53	53	53	53
54	54	54	54	54	54
55	55	55	55	55	55
56	56	56	56	56	56
57	57	57	57	57	57
58	58	58	58	58	58
59	59	59	59	59	59
60	60	60	60	60	60
61	61	61	61	61	61
62	62	62	62	62	62
63	63	63	63	63	63
64	64	64	64	64	64
65	65	65	65	65	65
66	66	66	66	66	66
67	67	67	67	67	67
68	68	68	68	68	68
69	69	69	69	69	69
70	70	70	70	70	70
71	71	71	71	71	71
72	72	72	72	72	72
73	73	73	73	73	73
74	74	74	74	74	74
75	75	75</			

Ash Glu Glu Val Met Phe Leu Thr Val Gln Val Lys Gly Pro Thr Gln

100 105 110

Glu Phe Lys Lys Arg Thr Thr Val Met Val Lys Asn Glu Asp Ser Leu

115	120	125
-----	-----	-----

Val	Phe	Val	Gln	Thr	Asp	Lys	Ser	Ile	Tyr	Lys	Pro	Gly	Gln	Thr	Val
130						135				140					
Lys	Phe	Arg	Val	Val	Ser	Met	Asp	Glu	Asn	Phe	His	Pro	Leu	Asn	Glu
145					150					155					160
Leu	Ile	Pro	Leu	Val	Tyr	Ile	Gln	Asp	Pro	Lys	Gly	Asn	Arg	Ile	Ala
				165					170					175	
Gln	Trp	Gln	Ser	Phe	Gln	Leu	Glu	Gly	Gly	Leu	Lys	Gln	Phe	Ser	Phe
			180					185					190		
Pro	Leu	Ser	Ser	Glu	Pro	Phe	Gln	Gly	Ser	Tyr	Lys	Val	Val	Val	Gln
		195				200						205			
Lys	Lys	Ser	Gly	Gly	Arg	Thr	Glu	His	Pro	Phe	Thr	Val	Glu	Glu	Phe
210					215						220				
Val	Leu	Pro	Lys	Phe	Glu	Val	Gln	Val	Thr	Val	Pro	Lys	Ile	Ile	Thr
225					230					235					240
Ile	Leu	Glu	Glu	Glu	Met	Asn	Val	Ser	Val	Cys	Gly	Leu	Tyr	Thr	Tyr
				245					250					255	
Gly	Lys	Pro	Val	Pro	Gly	His	Val	Thr	Val	Ser	Ile	Cys	Arg	Lys	Tyr
			260					265					270		
Ser	Asp	Ala	Ser	Asp	Cys	His	Gly	Glu	Asp	Ser	Gln	Ala	Phe	Cys	Glu
		275					280					285			
Lys	Phe	Ser	Gly	Gln	Leu	Asn	Ser	His	Gly	Cys	Phe	Tyr	Gln	Gln	Val
290					295						300				
Lys	Thr	Lys	Val	Phe	Gln	Leu	Lys	Arg	Lys	Glu	Tyr	Glu	Met	Lys	Leu
305					310					315					320
His	Thr	Glu	Ala	Gln	Ile	Gln	Glu	Glu	Gly	Thr	Val	Val	Glu	Leu	Thr
				325					330					335	
Gly	Arg	Gln	Ser	Ser	Glu	Ile	Thr	Arg	Thr	Ile	Thr	Lys	Leu	Ser	Phe
			340					345					350		
Val	Lys	Val	Asp	Ser	His	Phe	Arg	Gln	Gly	Ile	Pro	Phe	Phe	Gly	Gln
		355					360					365			
Val	Arg	Leu	Val	Asp	Gly	Lys	Gly	Val	Pro	Ile	Pro	Asn	Lys	Val	Ile
370					375						380				
Phe	Ile	Arg	Gly	Asn	Glu	Ala	Asn	Tyr	Tyr	Ser	Asn	Ala	Thr	Thr	Asp
385					390					395					400
Glu	His	Gly	Leu	Val	Gln	Phe	Ser	Ile	Asn	Thr	Thr	Asn	Val	Met	Gly
				405					410					415	
Thr	Ser	Leu	Thr	Val	Arg	Val	Asn	Tyr	Lys	Asp	Arg	Ser	Pro	Cys	Tyr
			420					425					430		
Gly	Tyr	Gln	Trp	Val	Ser	Glu	Glu	His	Glu	Glu	Ala	His	His	Thr	Ala
		435				440					445				
Tyr	Leu	Val	Phe	Ser	Pro	Ser	Lys	Ser	Phe	Val	His	Leu	Glu	Pro	Met
450					455						460				
Ser	His	Glu	Leu	Pro	Cys	Gly	His	Thr	Gln	Thr	Val	Gln	Ala	His	Tyr
465					470					475					480
Ile	Leu	Asn	Gly	Gly	Thr	Leu	Leu	Gly	Leu	Lys	Lys	Leu	Ser	Phe	Tyr
				485				490						495	
Tyr	Leu	Ile	Met	Ala	Lys	Gly	Gly	Ile	Val	Arg	Thr	Gly	Thr	His	Gly
			500					505					510		
Leu	Leu	Val	Lys	Gln	Glu	Asp	Met	Lys	Gly	His	Phe	Ser	Ile	Ser	Ile
		515				520					525				
Pro	Val	Lys	Ser	Asp	Ile	Ala	Pro	Val	Ala	Arg	Leu	Leu	Ile	Tyr	Ala
					535						540				
Val	Leu	Pro	Thr	Gly	Asp	Val	Ile	Gly	Asp	Ser	Ala	Lys	Tyr	Asp	Val
545					550					555					560
Glu	Asn	Cys	Leu	Ala	Asn	Lys	Val	Asp	Leu	Ser	Phe	Ser	Pro	Ser	Gln
				565				570						575	
Ser	Leu	Pro	Ala	Ser	His	Ala	His	Leu	Arg	Val	Thr	Ala	Ala	Pro	Gln
			580					585					590		

Ser	Val	Cys	Ala	Leu	Arg	Ala	Val	Asp	Gln	Ser	Val	Leu	Leu	Met	Lys
		595					600					605			
Pro	Asp	Ala	Glu	Leu	Ser	Ala	Ser	Ser	Val	Tyr	Asn	Leu	Leu	Pro	Glu
	610					615					620				
Lys	Asp	Leu	Thr	Gly	Phe	Pro	Gly	Pro	Leu	Asn	Asp	Gln	Asp	Asp	Glu
625					630					635					640
Asp	Cys	Ile	Asn	Arg	His	Asn	Val	Tyr	Ile	Asn	Gly	Ile	Thr	Tyr	Thr
				645					650					655	
Pro	Val	Ser	Ser	Thr	Asn	Glu	Lys	Asp	Met	Tyr	Ser	Phe	Leu	Glu	Asp
			660					665					670		
Met	Gly	Leu	Lys	Ala	Phe	Thr	Asn	Ser	Lys	Ile	Arg	Lys	Pro	Lys	Met
		675					680					685			
Cys	Pro	Gln	Leu	Gln	Gln	Tyr	Glu	Met	His	Gly	Pro	Glu	Gly	Leu	Arg
	690					695					700				
Val	Gly	Phe	Tyr	Glu	Ser	Asp	Val	Met	Gly	Arg	Gly	His	Ala	Arg	Leu
705					710					715					720
Val	His	Val	Glu	Glu	Pro	His	Thr	Glu	Thr	Val	Arg	Lys	Tyr	Phe	Pro
				725					730					735	
Glu	Thr	Trp	Ile	Trp	Asp	Leu	Val	Val	Val	Asn	Ser	Ala	Gly	Val	Ala
			740					745					750		
Glu	Val	Gly	Val	Thr	Val	Pro	Asp	Thr	Ile	Thr	Glu	Trp	Lys	Ala	Gly
		755					760					765			
Ala	Phe	Cys	Leu	Ser	Glu	Asp	Ala	Gly	Leu	Gly	Ile	Ser	Ser	Thr	Ala
	770					775					780				
Ser	Leu	Arg	Ala	Phe	Gln	Pro	Phe	Phe	Val	Glu	Leu	Thr	Met	Pro	Tyr
785					790					795					800
Ser	Val	Ile	Arg	Gly	Glu	Ala	Phe	Thr	Leu	Lys	Ala	Thr	Val	Leu	Asn
				805					810					815	
Tyr	Leu	Pro	Lys	Cys	Ile	Arg	Val	Ser	Val	Gln	Leu	Glu	Ala	Ser	Pro
			820					825					830		
Ala	Phe	Leu	Ala	Val	Pro	Val	Glu	Lys	Glu	Gln	Ala	Pro	His	Cys	Ile
		835					840					845			
Cys	Ala	Asn	Gly	Arg	Gln	Thr	Val	Ser	Trp	Ala	Val	Thr	Pro	Lys	Ser
	850					855					860				
Leu	Gly	Asn	Val	Asn	Phe	Thr	Val	Ser	Ala	Glu	Ala	Leu	Glu	Ser	Gln
865					870					875					880
Glu	Leu	Cys	Gly	Thr	Glu	Val	Pro	Ser	Val	Pro	Glu	His	Gly	Arg	Lys
				885					890					895	
Asp	Thr	Val	Ile	Lys	Pro	Leu	Leu	Val	Glu	Pro	Glu	Gly	Leu	Glu	Lys
			900					905					910		
Glu	Thr	Thr	Phe	Asn	Ser	Leu	Leu	Cys	Pro	Ser	Gly	Gly	Glu	Val	Ser
		915					920					925			
Glu	Glu	Leu	Ser	Leu	Lys	Leu	Pro	Pro	Asn	Val	Val	Glu	Glu	Ser	Ala
					935						940				
Arg	Ala	Ser	Val	Ser											

Phe Ile Asp Glu Ala His Ile Thr Gln Ala Leu Ile Trp Leu Ser Gln
1060 1065 1070
Arg Gln Lys Asp Asn Gly Cys Phe Arg Ser Ser Gly Ser Leu Leu Asn
1075 1080 1085
Asn Ala Ile Lys Gly Gly Val Glu Asp Glu Val Thr Leu Ser Ala Tyr
1090 1095 1100
Ile Thr Ile Ala Leu Leu Glu Ile Pro Leu Thr Val Thr His Pro Val
1105 1110 1115 1120
Val Arg Asn Ala Leu Phe Cys Leu Glu Ser Ala Trp Lys Thr Ala Gln
1125 1130 1135
Glu Gly Asp His Gly Ser His Val Tyr Thr Lys Ala Leu Leu Ala Tyr
1140 1145 1150
Ala Phe Ala Leu Ala Gly Asn Gln Asp Lys Arg Lys Glu Val Leu Lys
1155 1160 1165
Ser Leu Asn Glu Glu Ala Val Lys Lys Asp Asn Ser Val His Trp Glu
1170 1175 1180
Arg Pro Gln Lys Pro Lys Ala Pro Val Gly His Phe Tyr Glu Pro Gln
1185 1190 1195 1200
Ala Pro Ser Ala Glu Val Glu Met Thr Ser Tyr Val Leu Leu Ala Tyr
1205 1210 1215
Leu Thr Ala Gln Pro Ala Pro Thr Ser Glu Asp Leu Thr Ser Ala Thr
1220 1225 1230
Asn Ile Val Lys Trp Ile Thr Lys Gln Gln Asn Ala Gln Gly Gly Phe
1235 1240 1245
Ser Ser Thr Gln Asp Thr Val Val Ala Leu His Ala Leu Ser Lys Tyr
1250 1255 1260
Gly Ala Ala Thr Phe Thr Arg Thr Gly Lys Ala Ala Gln Val Thr Ile
1265 1270 1275 1280
Gln Ser Ser Gly Thr Phe Ser Ser Lys Phe Gln Val Asp Asn Asn Asn
1285 1290 1295
Arg Leu Leu Leu Gln Gln Val Ser Leu Pro Glu Leu Pro Gly Glu Tyr
1300 1305 1310
Ser Met Lys Val Thr Gly Glu Gly Cys Val Tyr Leu Gln Thr Ser Leu
1315 1320 1325
Lys Tyr Asn Ile Leu Pro Glu Lys Glu Glu Phe Pro Phe Ala Leu Gly
1330 1335 1340
Val Gln Thr Leu Pro Gln Thr Cys Asp Glu Pro Lys Ala His Thr Ser
1345 1350 1355 1360
Phe Gln Ile Ser Leu Ser Val Ser Tyr Thr Gly Ser Arg Ser Ala Ser
1365 1370 1375
Asn Met Ala Ile Val Asp Val Lys Met Val Ser Gly Phe Ile Pro Leu
1380 1385 1390
Lys Pro Thr Val Lys Met Leu Glu Arg Ser Asn His Val Ser Arg Thr
1395 1400 1405
Glu Val Ser Ser Asn His Val Leu Ile Tyr Leu Asp Lys Val Ser Asn
1410 1415 1420
Gln Thr Leu Ser Leu Phe Phe Thr Val Leu Gln Asp Val Pro Val Arg
1425 1430 1435 1440
Asp Leu Lys Pro Ala Ile Val Lys Val Tyr Asp Tyr Tyr Glu Thr Asp
1445 1450 1455
Glu Phe Ala Ile Ala Glu Tyr Asn Ala Pro Cys Ser Lys Asp Leu Gly
1460 1465 1470
Asn Ala

<210> 6
<211> 14896
<212> DNA

<213> Homo sapiens

<400> 6

cagcgggtgcg	agctccaggg	ccatgcactg	aggaggcgga	aacaagggga	gccccagag	60
ctccatcaag	ccccctccaa	aggctcccct	acccgggtcca	cgccccccac	ccccctccc	120
cgctcctcc	caattgtgca	tttttgcagc	cggaggcggc	tccgagatgg	ggctgtgagc	180
ttcgcccggg	gagggggaaa	gagcagcgag	gagtgaagcg	gggggggtggg	gtgaagggtt	240
tggatttcgg	ggcagggggc	gcacccccgt	cagcaggccc	tccccagggg	gctcggaact	300
ctacctcttc	accacgccc	ctggtgcgct	ttgccgaagg	aaagaataag	aacagagaag	360
gaggaggggg	aaaggaggaa	aagggggacc	ccccaaactgg	gggggggtgaa	ggagagaagt	420
agcaggacca	gaggggaagg	ggctgctgct	tgcctcagcc	cacaccatgc	tgaccccgcc	480
gttgctcctg	ctgctgcccc	tgtctcagc	tctgggtcgcg	gcggctatcg	acgcccctaa	540
gacttgcagc	cccaagcagt	ttgcctgcag	agatcaaata	acctgtatct	caaagggtctg	600
gcgggtgcgac	ggtgagaggg	actgcccaga	cggatctgac	gaggccccctg	agatttgtcc	660
acagagtaag	gcccagcgat	gccagccaaa	cgagcataac	tgcctgggta	ctgagctgtg	720
tgttcccatg	tcccgctct	gcaatggggg	ccaggactgc	atggacggct	cagatgaggg	780
gccccactgc	cgagagctcc	aaggcaactg	ctctcgctg	ggctgccagc	accatttgtgt	840
ccccacactc	gatgggccc	cctgctactg	caacagcagc	tttcagcttc	aggcagatgg	900
caagacctgc	aaagattttg	atgagtgtct	agtgtacggc	acctgcagcc	agctatgcac	960
caacacagac	ggctccttca	tatgtggctg	tgttgaagga	tacctcctgc	agccggataa	1020
ccgctcctgc	aaggccaaga	acgagccagt	agaccggccc	cctgtgctgt	tgatagccaa	1080
ctcccagaac	atcttggcca	cgtacctgag	tggggcccag	gtgtctacca	tcacacctac	1140
gagcacgcgg	cagaccacag	ccatggactt	cagctatgcc	aacgagaccg	tatgctgggt	1200
gcatgtttggg	gacagtgtct	ctcagacgca	gctcaagtgt	gcccgcctgc	ctggcctaaa	1260
gggcttcgtg	gatgagcaca	ccatcaacat	ctccctcagt	ctgcaccacg	tggaacagat	1320
ggccatcgac	tggctgacag	gcaacttcta	ctttgtggat	gacatcgatg	ataggatctt	1380
tgtctgcaac	agaaatgggg	acacatgtgt	cacattgcta	gacctggaac	tctacaacct	1440
caagggcatt	gccctggacc	ctgccatggg	gaagggtgtt	ttcactgact	atgggcagat	1500
cccaaagggtg	gaacgctgtg	acatggatgg	gcagaaccgc	accaagctcg	tcgacagcaa	1560
gattgtgttt	cctcatggca	tcacgctgga	cctggctcagc	cgcttgtct	actgggcaga	1620
tgcctatctg	gactatattg	aagtgggtgga	ctatgagggc	aaggggccgc	agaccatcat	1680
ccaggggcatc	ctgattgagc	acctgtacgg	cctgactgtg	tttgagaatt	atctctatgc	1740
caccaactcg	gacaatgcca	atgcccagca	gaagacgagt	gtgatccgtg	tgaaccgctt	1800
taacagcacc	gagtaccagg	ttgtcaccgg	ggtggacaag	ggtgggtgcc	tccacatcta	1860
ccaccagagg	cgtcagcccc	gagtgaggag	ccatgcctgt	gaaaacgacc	agtatgggaa	1920
gcccgggtggc	tgtcttgaca	tctgcctgct	ggccaacagc	cacaaggcgc	ggacctgccg	1980
ctgccgttcc	ggcttcagcc	tgggcagtg	cgggaagtca	tgcaagaagc	cggagcatga	2040
gctgttcctc	gtgtatggca	agggccggcc	aggcatcatc	cggggcatgg	atatgggggc	2100
caagggtccc	gatgagcaca	tgatccccat	tgaaaacctc	atgaaccccc	gagccctgga	2160
cttcacgct	gagaccggct	tcacttactt	tgccgacacc	accagctacc	tcattggccg	2220
ccagaagatt	gatggcactg	agcgggagac	catcctgaag	gacggcatcc	acaatgtgga	2280
gggtgtggcc	gtggactgga	tgggagacaa	tctgtactgg	acggacgatg	ggcccaaaaa	2340
gacaatcagc	gtggccaggg	tggagaaagc	tgtcagacc	cgcaagactt	taatcgaggg	2400
caaaatgaca	caccccaggg	ctattgtggg	ggatccactc	aatgggtgga	tgtactggac	2460
agactgggag	gaggacccca	aggacagtgc	gcgtggggcg	ctggagaggg	cgtggatgga	2520
tggctcacac	cgagacatct	ttgtcacctc	caagacagtg	ctttggccca	atgggctaag	2580
cctggacatc	ccggctgggc	gcctctactg	ggtggatgcc	ttctacgacc	gcatcgagac	2640
gatactgctc	aatggcacag	accggaagat	tgtgtatgaa	ggtcctgagc	tgaaccacgc	2700
ctttggcctg	tgtcaccatg	gcaactacct	cttctggact	gagtatcgga	gtggcagtgt	2760
ctaccgcttg	gaacgggggtg	taggaggcgc	acccccact	gtgaccttc	tgcgcagtga	2820
gcgccccccc	atctttgaga	tccgaatgta	tgatgccag	cagcagcaag	ttggcaccaa	2880
caaatgccgg	gtgaacaatg	gcggctgcag	cagcctgtgc	ttggccaccc	ctgggagccg	2940
ccagtgcgcc	tgtgctgagg	accaggtgtt	ggacgcagac	ggcgtcactt	gcttggcgaa	3000
cccatcctac	gtgcctccac	cccagtgcc	gccaggcgag	tttgctgtg	ccaacagccg	3060
ctgcatccag	gagcgttgga	agtgtgacgg	agacaacgat	tgcctggaca	acagtgatga	3120
ggccccagcc	ctctgccatc	agcacacctg	cccctcgga	cgattcaagt	gcgagaacaa	3180
ccgggtgcac	cccaaccgct	ggctctgcga	cggggacaat	gactgtggga	acagtgaaga	3240
tgagtccaat	gccacttggt	cagccccgac	ctgccccccc	aaccagttct	cctgtgccag	3300

CC5092-0454

tggccgctgc	atccccatct	cctggacgtg	tgatctggat	gacgactgtg	gggaccgctc	3360
tgatgagtct	gcttcgtgtg	cctatcccac	ctgcttcccc	ctgactcagt	ttacctgcaa	3420
caatggcaga	tgtatcaaca	tcaactggag	atgcgacaat	gacaatgact	gtggggacaa	3480
cagtgcgaa	gccggctgca	gccactcctg	ttctagcacc	cagttcaagt	gcaacagcgg	3540
gcgttgcac	cccgagcact	ggacctgcga	tggggacaat	gactgcggag	actacagtga	3600
tgagacacac	gccaaactgca	ccaaccaggc	cacgaggccc	cctgggtggct	gccacactga	3660
tgagttccag	tgccggctgg	atggactatg	catccccctg	cgggtggcgct	gcgatgggga	3720
cactgactgc	atggactcca	gcgatgagaa	gagctgtgag	ggagtgaccc	acgtctgcga	3780
tcccagtgtc	aagtttggct	gcaaggactc	agctcgggtg	atcagcaaag	cgtgggtgtg	3840
tgatggcgac	aatgactgtg	aggataactc	ggacgaggag	aactgcgagt	ccctggcctg	3900
caggccaccc	tcgcaccctt	gtgccaaaca	cacctcagtc	tgcctgcccc	ctgacaagct	3960
gtgtgatggc	aacgacgact	gtggcgacgg	ctcagatgag	ggcgagctct	gcgaccagtg	4020
ctctctgaat	aacgggtggct	gcagccacaa	ctgctcagtg	gcacctggcg	aaggcattgt	4080
gtgttcctgc	cctctggggca	tggagctggg	gcccgcacaac	cacacctgcc	agatccagag	4140
ctactgtgcc	aagcatctca	aatgcagcca	aaagtgcgac	cagaacaagt	tcagcgtgaa	4200
gtgctcctgc	tacgagggct	gggtcctgga	acctgacggc	gagagctgcc	gcagcctgga	4260
ccccttcaag	ccgttcatca	ttttctccaa	ccgccatgaa	atccggcgca	tcgatcttca	4320
caaaggagac	tacagcgtcc	tggtgcccg	cctgcgcaac	accatcgccc	tggacttcca	4380
cctcagccag	agcgccctct	actggaccga	cgtgggtggag	gacaagatct	accgcgggaa	4440
gctgctggac	aacggagccc	tgactagttt	cgagggtggg	attcagtatg	gcctggccac	4500
acccgagggc	ctggctgtag	actggattgc	aggcaacatc	tactgggtgg	agagtaacct	4560
ggatcagatc	gaggtggcca	agctggatgg	gacctccgg	accacctgc	tggccggtga	4620
cattgagcac	ccaagggcaa	tcgcactgga	tccccgggat	gggatcctgt	tttggacaga	4680
ctgggatgcc	agcctgcccc	gcattgaggc	agcctccatg	agtggggctg	ggcgccgcac	4740
cgtgcaccgg	gagaccggct	ctgggggctg	gcccacggg	ctcaccgtgg	actacctgga	4800
gaagcgcac	ctttggattg	acgccaggtc	agatgccatt	tactcagccc	gttacgacgg	4860
ctctggccac	atggaggtgc	ttcgggggaca	cgagttcctg	tcgcaccctg	ttgcagtgac	4920
gctgtacggg	ggggaggtct	actggactga	ctggcgaaaca	aacacactgg	ctaaggccaa	4980
caagtggacc	ggccacaatg	tcaccgtggg	acagaggacc	aacacccagc	cctttgacct	5040
gcaggtgtac	caccctccc	gccagcccat	ggctcccaat	ccctgtgagg	ccaatggggg	5100
ccagggcccc	tgctcccacc	tgtgtctcat	caactacaa	cggaccgtgt	cctgcgcctg	5160
ccccacctc	atgaagctcc	acaaggacaa	caccacctgc	tatgagttta	agaagttcct	5220
gctgtacgca	cgtcagatgg	agatccgagg	tgtggacctg	gatgtccct	actacaacta	5280
catcatctcc	ttcacgggtgc	ccgacatcga	caacgtcaca	gtgctagact	acgatgcccg	5340
cgagcagcgt	gtgtactggg	ctgacgtgcg	gacacaggcc	atcaagcggg	ccttcatcaa	5400
cggcacaggc	gtggagacag	tcgtctctgc	agacttgcca	aatgcccacg	ggctggctgt	5460
ggactgggtc	tcccgaacc	tgttctggac	aagctatgac	accaataaga	agcagatcaa	5520
tgtggcccg	ctggatggct	ccttcaagaa	cgcagtgggtg	cagggcctgg	agcagcccca	5580
tggccttgtc	gtccaccctc	tgcgtgggaa	gctctactgg	accgatgggtg	acaacatcag	5640
catggccaac	atggatggca	gcaatcgcac	cctgctcttc	agtggccaga	agggccccgt	5700
gggcctggct	attgacttcc	ctgaaagcaa	actctactgg	atcagctccg	ggaaccatac	5760
catcaaccgc	tgcaacctgg	atgggagtgg	gctggagggtc	atcgatgcca	tgcggagcca	5820
gctgggcaag	gccaccgccc	tggccatcat	gggggacaa	ctgtggtggg	ctgatcaggt	5880
gtcggaaaag	atgggcacat	gcagcaaggc	tgacggctcg	ggctccgtgg	tccttcggaa	5940
cagcaccacc	ctgggtgatgc	acatgaagg	ctatgacgag	agcatccagc	tggaccataa	6000
gggcaccaac	ccctgcagtg	tcaacaacgg	tgactgctcc	cagctctgcc	tgcccacgtc	6060
agagacgacc	cgtccctgca	tgtgcacagc	cggctatagc	ctccggagtg	gccagcaggc	6120
ctgcgagggc	gtaggttcct	ttctcctgta	ctctgtgcat	gagggaatca	ggggaattcc	6180
cctggatccc	aatgacaagt	cagatgccct	ggtcccagt	tccgggacct	cgtggctgt	6240
cggcatcgac	ttccacgctg	aaaatgacac	catctactgg	gtggacatgg	gcctgagcac	6300
gatcagccgg	gccaaagcggg	accagacgtg	gcgtgaagac	gtggtgacca	atggcattgg	6360
ccgtgtggag	ggcattgcag	tggactggat	cgcaggcaac	atctactgga	cagaccaggg	6420
ctttgatgtc	atcgaggctg	cccggctcaa	tggctccttc	cgtacgtgg	tgatctccca	6480
gggtctagac	aagccccggg	ccatcacctg	ccaccgggag	aaagggtact	tgttctggac	6540
tgagtgggg	cagtatccgc	gtattgagcg	gtctcggcta	gatggcacgg	agcgtgtgg	6600
gctgggtcaac	gtcagcatca	gctggcccaa	cggcatctca	gtggactacc	aggatgggaa	6660
gctgtactgg	tcgatgcac	ggacagacaa	gattgaacgg	atcgacctgg	agacaggtga	6720
gaaccgcgag	gtggttctgt					

ggatttcatc	tactggagtg	acaggactca	tgccaacggc	tctatcaagc	gcgggagcaa	6840
agacaatgcc	acagactccg	tgcccctgcg	aaccggcatc	ggcgtccagc	ttaaagacat	6900
caaagtcttc	aaccgggacc	ggcagaaagg	caccaacgtg	tgcgcggttg	ccaatggcgg	6960
gtgccagcag	ctgtgcctgt	accggggccg	tgggcagcgg	gcctgcgcct	gtgcccacgg	7020
gatgctggct	gaagacggag	catcgtgccg	cgagtatgcc	ggctacctgc	tctactcaga	7080
gcgcaccatt	ctcaagagta	tccacctgtc	ggatgagcgc	aacctcaatg	cgcccgtgca	7140
gcccttcgag	gaccttgagc	acatgaagaa	cgtcatcgcc	ctggcctttg	actaccgggc	7200
aggcacctct	ccgggcaccc	ccaatcgc at	cttcttcagc	gacatccact	ttgggaacat	7260
ccaacagatc	aacgacgatg	gctccaggag	gatcaccatt	gtggaaaacg	tgggctccgt	7320
ggaaggcctg	gcctatcacc	gtggctggga	cactctctat	tggacaagct	acacgacatc	7380
caccatcacg	cgccacacag	tggaccagac	ccgcccaggg	gccttcgagc	gtgagaccgt	7440
catcactatg	tctggagatg	accacccacg	ggccttcggt	ttggacgagt	gccagaacct	7500
catgtttctg	accaactgga	atgagcagca	tcccagcatc	atgcggggcg	cgctctcggg	7560
agccaatgtc	ctgaccctta	tcgagaagga	catccgtacc	cccaatggcc	tggccatcga	7620
ccaccgtgcc	gagaagctct	acttctctga	cgccaccctg	gacaagatcg	agcggtgcca	7680
gtatgacggc	tcccaccgct	atgtgatcct	aaagtccagag	cctgtccacc	ccttcgggct	7740
ggcctgtgat	ggggagcaca	ttttctggac	tgactgggtg	cggcggggcag	tgcagcgggc	7800
caacaagcac	gtgggcagca	acatgaagct	gctgcgcgtg	gacatcccc	agcagcccat	7860
gggcatcatc	gccgtggcca	acgacaccaa	cagctgtgaa	ctctctccat	gccgaatcaa	7920
caacgggtgg	tgccaggacc	tgtgtctgct	cactcaccag	ggccatgtca	actgctcatg	7980
ccgagggggc	cgaatcctcc	aggatgacct	cacctgccga	gcggtgaatt	cctcttgccg	8040
agcacaagat	gagtttgagt	gtgccaatgg	cgagtgcac	aacttcagcc	tgacctgcca	8100
cggcgtcccc	cactgcaagg	acaagtcgga	tgagaagcca	tcctactgca	actcccgcgc	8160
ctgcaagaag	actttccggc	agtgcagcaa	tgggcgctgt	gtgtccaaca	tgctgtggtg	8220
caacggggcc	gacgactgtg	gggatggctc	tgacgagatc	ccttgcaaca	agacagcctg	8280
tgggtgtggg	gagttccgct	gccgggacgg	gacctgcac	gggaactcca	gccgctgcaa	8340
ccagtttgtg	gattgtgagg	acgcctcaga	tgagatgaac	tgcagtgcca	ccgactgcag	8400
cagctacttc	cgctggggcg	tgaagggcgt	gctcttccag	ccctgcgagc	ggacctcact	8460
ctgctacgca	cccagctggg	tgtgtgatgg	cgccaatgac	tgtggggact	acagtgatga	8520
gcgcgactgc	ccaggtgtga	aacgccccag	atgcctctg	aattacttcg	cctgccctag	8580
tgggcgctgc	atccccatga	gctggacgtg	tgacaaagag	gatgactgtg	aacatggcca	8640
ggacgagacc	cactgcaaca	agttctgctc	agaggcccag	tttgagtgcc	agaaccatcg	8700
ctgcatctcc	aagcagtggc	tgtgtgacgg	cagcgatgac	tgtggggatg	gctcagacga	8760
ggctgctcac	tgtgaaggca	agacgtgcgg	cccctcctcc	ttctcctgcc	ctggcaccca	8820
cgtgtgcgtc	cccgagcgct	ggctctgtga	cggtgacaaa	gactgtgctg	atggtgcaga	8880
cgagagcatc	gcagctgggt	gcttgtacaa	cagcacttgt	gacgaccgtg	agttcatgtg	8940
ccagaaccgc	cagtgcaccc	ccaagcactt	cgtgtgtgac	cacgaccgtg	actgtgcaga	9000
tggctctgat	gagtcccccg	agtgtgagta	cccgaacctg	ggccccagtg	agttccgctg	9060
tgccaatggg	cgtgtcttga	gctcccgcga	gtgggagtgt	gatggcgaga	atgactgcca	9120
cgaccagagt	gacgaggctc	ccaagaaccc	acactgcacc	agcccagagc	acaagtgcaa	9180
tgctcgtca	cagttcctgt	gcagcagtgg	gcgctgtgtg	gctgaggcac	tgctctgcaa	9240
cggccaggat	gactgtggcg	acagctcgga	cgagcgtggc	tgccacatca	atgagtgtct	9300
cagccgcaag	ctcagtggct	gcagccagga	ctgtgaggac	ctcaagatcg	gcttcaagtg	9360
ccgctgtcgc	cctggcttcc	ggctgaagga	tgacggccgg	acgtgtgctg	atgtggacga	9420
gtgcagcacc	accttcccct	gcagccagcg	ctgcatcaac	acccatggca	gctataagtg	9480
tctgtgtgtg	gagggctatg	caccccgcgg	cggcgacccc	cacagctgca	aggctgtgac	9540
tgacgaggaa	ccgtttctga	tcttcgccaa	ccggtactac	ctgcgcaagc	tcaacctgga	9600
cgggtccaac	tacacgttac	ttaagcaggg	cctgaacaac	gccgttgctt	tggattttga	9660
ctaccgagag	cagatgatct	actggacaga	tgtgaccacc	cagggcagca	tgatccgaag	9720
gatgcacctt	aacgggagca	atgtgcaggt	cctacaccgt	acaggcctca	gcaaccccca	9780
tgggctggct	gtggactggg	tgggtggcaa	cctgtactgg	tgcgacaaa	gccgggacac	9840
catcgaggtg	tccaagctca	atggggccta	tgggacgggt	ctggctagct	ctggcctccg	9900
tgagcccagg	gctctgggtg	tggatgtgca	gaatgggtac	ctgtactgga	cagactgggg	9960
tgaccattca	ctgatcggcc	gcacgcgcac	ggatgggtcc	agccgcagcg	tcacgttgga	10020
caccaagatc	acatggccca	atggcctgac	gctggactat	gtcactgagc	gcactctactg	10080
ggccgacgcc	cgcgaggact	acattgaatt	tgccagcctg	gatgggtcca	atcgccacgt	10140
tgtgctgagc	caggacatcc	cgcacatctt	tgcactgacc	ctgtttgagg	actacgtcta	10200

aacgctcctc	atcagcacgc	tgcaccggcc	catggacctg	catgtcttcc	atgccctgcg	10320
ccagccagac	gtgccaatc	acccctgcaa	ggtcaacaat	ggtggctgca	gcaacctgtg	10380
cctgctgtcc	cccgggggag	ggcacaatg	tgcctgcccc	accaacttct	acctgggcag	10440
cgatgggcgc	acctgtgtgt	ccaactgcac	ggctagccag	tttgtatgca	agaacgacaa	10500
gtgcatcccc	ttctgggtga	agtgtgacac	cgaggacgac	tgcggggacc	actcagacga	10560
gcccccgga	tgccctgagt	tcaagtgccg	gccccggacag	ttccagtgtc	ccacaggtat	10620
ctgcacaaac	cctgccttca	tctgcgatgg	cgacaatgac	tgccaggaca	acagtgcgca	10680
ggccaactgt	gacatccacg	tctgcttgcc	cagtgcgttc	aaatgcacca	acaccaaccg	10740
ctgtattccc	ggcatcttcc	gctgcaatgg	gcaggacaac	tgcggagatg	gggaggatga	10800
gagggactgc	cccagaggtga	cctgcgcccc	caaccagttc	cagtgtctca	ttaccaaacg	10860
gtgcatcccc	cgggtctggg	tctgcgaccg	ggacaatgac	tgtgtggatg	gcagtgcgca	10920
gccccgcaac	tgcacccaga	tgacctgtgg	tgtggacgag	ttccgctgca	aggattcggg	10980
ccgctgcac	ccagcgcgtt	ggaagtgtga	cggagaggat	gactgtgggg	atggctcgga	11040
tgagcccaag	gaagagtgtg	atgaacgcac	ctgtgagcca	taccagtctc	gctgcaagaa	11100
caaccgctgc	gtgcccggcc	gctggcagtg	cgactacgac	aacgattgcg	gtgacaactc	11160
cgatgaagag	agctgcaccc	ctcggccctg	ctccgagagt	gagttctcct	gtgccaacgg	11220
ccgctgcac	gcggggcgct	ggaaatgcga	tggagaccac	gactgcgcgg	acggctcgga	11280
cgagaaagac	tgcaccccc	gctgtgacat	ggaccagttc	cagtgcgaaga	gcggccactg	11340
catccccctg	cgttggcgct	gtgacgcaga	cggcgactgc	atggacggca	gcgacgagga	11400
ggcctgcggc	actggcgtgc	ggacctgccc	cctggacgag	ttccagtgc	acaacacctt	11460
gtgcaagccg	ctggcctgga	agtgcgatgg	cgaggatgac	tgtggggaca	actcagatga	11520
gaaccccgag	gagtgtgccc	ggttcgtgtg	ccctcccaac	cggcccttcc	gttgcaagaa	11580
tgaccgcgtc	tgtctgtgga	tcggggcgcca	atgcatgggc	acggacaact	gtggggatgg	11640
gactgatgaa	gaggactgtg	agccccccac	agccacacac	acccactgca	aagacaagaa	11700
ggagtcttctg	tgccggaacc	agcgtgcctt	ctcctcctcc	ctgcgctgca	acatgttcga	11760
tgactgcggg	gacggctctg	acgaggagga	ctgcagcatc	gaccccaagc	tgaccagctg	11820
cgccaccaat	gccagcatct	gtggggacga	ggcacgctgc	gtgcgcaccg	agaaagcggc	11880
ctactgtgcc	tgccgctcgg	gcttccacac	cgtgcccggc	cagcccggat	gccaagacat	11940
caacgagtgc	ctgcgcttcg	gcacctgtct	ccagctctgc	aacaacacca	agggcggcca	12000
cctctgcagc	tgcgctcgga	acttcatgaa	gacgcacaac	acctgcaagg	ccgaaggctc	12060
tgagtaccag	gtcctgtaca	tcgctgatga	caatgagatc	cgcagcctgt	tccccggcca	12120
cccccatctg	gcttacgagc	aggcattcca	gggtgacgag	agtgtccgca	ttgatgctat	12180
ggatgtccat	gtcaaggctg	gccgtgtcta	ttggaccaac	tggcacacgg	gcaccatctc	12240
ctaccgcagc	ctgccacctg	ctgcgcctcc	taccacttcc	aaccgccacc	ggcgacagat	12300
tgaccggggg	gtcaccaccc	tcaacatttc	agggctgaag	atgccccagag	gcatcgccat	12360
cgactgggtg	gccggaaacg	tgtactggac	cgactcgggc	cgagatgtga	ttgaggtggc	12420
gcagatgaag	ggcgagaacc	gcaagacgct	catctcgggc	atgattgacg	agccccacgc	12480
cattgtgggtg	gacccactga	gggggaccat	gtactggtca	gactggggca	accaccccaa	12540
gattgagacg	gcagcgatgg	atgggacgct	tcgggagaca	ctggtgcagg	acaacattca	12600
gtggcccaca	ggcctggccg	tggattatca	caatgagcgg	ctgtactggg	cagacgccaa	12660
gctttcagtc	atcggcagca	tccggctcaa	tggcacggac	ccattgtggg	ctgctgacag	12720
caaacgaggg	ctaagtccac	ccttcagcat	cgacgtcttt	gaggattaca	tctatgggtg	12780
cacctacatc	aataatcgtg	tcttcaagat	ccataagttt	ggccacagcc	ccttgggtcaa	12840
cctgacaggg	ggcctgagcc	acgcctctga	cgtggctcct	taccatcagc	acaagcagcc	12900
cgaagtgacc	aacccatgtg	accgcaagaa	atgcgagtgg	ctctgcctgc	tgagccccag	12960
tgggcctgtc	tgcacctgtc	ccaatgggaa	gcggctggac	aacggcacat	gcgtgcctgt	13020
gccctctcca	acgccccccc	cagatgctcc	ccggcctgga	acctgtaacc	tgcagtgtct	13080
caacgggtgg	agctgtttcc	tcaatgcacg	gaggcagccc	aagtgccgct	gccaaccccc	13140
ctacacgggt	gacaagtgtg	aactggacca	gtgctgggag	cactgtcgca	atggggggcac	13200
ctgtgctgcc	tccccctctg	gcatgcccac	gtgccgggtg	cccacgggct	tcacggggccc	13260
caaatgcacc	cagcaggtgt	gtgcgggcta	ctgtgccaac	aacagcacct	gcactgtcaa	13320
ccagggcaac	cagccccagt	gccgatgcct	acccggcttc	ctggggcgacc	gctgccagta	13380
ccggcagtgc	tctggctact	gtgagaactt	tggcacatgc	cagatggctg	ctgatggctc	13440
ccgacaatgc	cgtgcactg	cctactttga	gggatcgagg	tgtgaggtga	acaagtgcag	13500
ccgctgtctc	gaagggggcct	gtgtgggtcaa	caagcagagt	ggggatgtca	cctgcaactg	13560
cacggatggc	cgggtggccc	ccagctgtct	gacctgcgtc	ggccactgca	gcaatggcgg	13620
ctcctgtacc	atgaacagca	aatgatgcc	tgagtgccag	tgcacacccc	acatgacagg	13680
gccccgggtg	gaggagcacg	tcttcagcca	gcagcagcca	ggacatatag	cctccatcct	13740

```

aatccctctg ctgttgctgc tgctgctggt tctgggtggcc ggagtgggtat tctgggtataa 13800
gcggcgagtc caaggggcta agggcttcca gcaccaacgg atgaccaacg gggccatgaa 13860
cgtggagatt ggaaacccca cctacaagat gtacgaaggc ggagagcctg atgatgtggg 13920
aggcctactg gacgctgact ttgccctgga ccctgacaag cccaccaact tcaccaaccc 13980
cgtgtatgcc acactctaca tgggggggcca tggcagtcgc cactccctgg ccagcacgga 14040
cgagaagcga gaactcctgg gccggggggccc tgaggacgag atagggggacc ccttgggcata 14100
gggccctgcc ccgtcggact gccccagaa agcctcctgc cccctgccgg tgaagtcctt 14160
cagtgaagccc ctccccagcc agcccttccc tggccccgcc ggatgtataa atgtaaaaat 14220
gaaggaatta cattttatat gtgagcgagc aagccggcaa gcgagcacag tattatttct 14280
ccatccccctc cctgcctgct ccttggcacc cccatgctgc cttcaggagg acaggcaggg 14340
agggcttggg gctgcacctc ctaccctccc accagaacgc accccactgg gagagctggt 14400
ggtgcagcct tccccctcct gtataagaca ctttgccaag gctctccccct ctcgccccat 14460
ccctgcttgc ccgctcccac agcttcctga gggctaattc tgggaaggga gagttctttg 14520
ctgccccctgt ctggaagacg tggctctggg tgaggtaggc gggaaaggat ggagtgtttt 14580
agttcttggg ggaggccacc ccaaacccca gcccacactc cagggggcacc tatgagatgg 14640
ccatgctcaa cccccctccc agacaggccc tccctgtctc caggggcccc accgaggttc 14700
ccagggtctg agacttcctc tggtaaacad tcctccagcc tccccctccc tggggacgcc 14760
aaggaggtgg gccacaccca ggaagggaaa gcgggcagcc ccgttttggg gacgtgaacg 14820
ttttaataat ttttgctgaa ttctttacaa ctaaataaca cagatattct tataaataaa 14880
attgtaaaaa aaaaaa 14896

```

```

<210> 7
<211> 126
<212> PRT
<213> Homo sapiens

```

```

<400> 7
Ile Ala Leu Asp Phe His Leu Ser Gln Ser Ala Leu Tyr Trp Thr Asp
 1          5          10          15
Val Val Glu Asp Lys Ile Tyr Arg Gly Lys Leu Leu Asp Asn Gly Ala
          20          25          30
Leu Thr Ser Phe Glu Val Val Ile Gln Tyr Gly Leu Ala Thr Pro Glu
          35          40          45
Gly Leu Ala Val Asp Trp Ile Ala Gly Asn Ile Tyr Trp Val Glu Ser
          50          55          60
Asn Leu Asp Gln Ile Glu Val Ala Lys Leu Asp Gly Thr Leu Arg Thr
65          70          75          80
Thr Leu Leu Ala Gly Asp Ile Glu His Pro Arg Ala Ile Ala Leu Asp
          85          90          95
Pro Arg Asp Gly Ile Leu Phe Trp Thr Asp Trp Asp Ala Ser Leu Pro
          100          105          110
Arg Ile Glu Ala Ala Ser Met Ser Gly Ala Gly Arg Arg Thr
          115          120          125

```

```

<210> 8
<211> 153
<212> PRT
<213> Homo sapiens

```

```

<400> 8
Leu Leu Gln Gln Val Ser Leu Pro Glu Leu Pro Gly Glu Tyr Ser Met
 1          5          10          15
Lys Val Thr Gly Glu Gly Cys Val Tyr Leu Gln Thr Ser Leu Lys Tyr
          20          25          30
Asn Ile Leu Pro Glu Lys Glu Glu Phe Pro Phe Ala Leu Gly Val Gln
          35          40          45

```

Thr	Leu	Pro	Gln	Thr	Cys	Asp	Glu	Pro	Lys	Ala	His	Thr	Ser	Phe	Gln
50						55					60				
Ile	Ser	Leu	Ser	Val	Ser	Tyr	Thr	Gly	Ser	Arg	Ser	Ala	Ser	Asn	Met
65					70					75					80
Ala	Ile	Val	Asp	Val	Lys	Met	Val	Ser	Gly	Phe	Ile	Pro	Leu	Lys	Pro
				85					90					95	
Thr	Val	Lys	Met	Leu	Glu	Arg	Ser	Asn	His	Val	Ser	Arg	Thr	Glu	Val
			100					105					110		
Ser	Ser	Asn	His	Val	Leu	Ile	Tyr	Leu	Asp	Lys	Val	Ser	Asn	Gln	Thr
		115					120					125			
Leu	Ser	Leu	Phe	Phe	Thr	Val	Leu	Gln	Asp	Val	Pro	Val	Arg	Asp	Leu
	130					135					140				
Lys	Pro	Ala	Ile	Val	Lys	Val	Tyr	Asp							
145					150										

<210> 9
 <211> 138
 <212> PRT
 <213> Homo sapiens

Met	Lys	Val	Thr	Gly	Glu	Gly	Cys	Val	Tyr	Leu	Gln	Thr	Ser	Leu	Lys
1				5					10					15	
Tyr	Asn	Ile	Leu	Pro	Glu	Lys	Glu	Glu	Phe	Pro	Phe	Ala	Leu	Gly	Val
			20					25					30		
Gln	Thr	Leu	Pro	Gln	Thr	Cys	Asp	Glu	Pro	Lys	Ala	His	Thr	Ser	Phe
		35					40					45			
Gln	Ile	Ser	Leu	Ser	Val	Ser	Tyr	Thr	Gly	Ser	Arg	Ser	Ala	Ser	Asn
	50					55					60				
Met	Ala	Ile	Val	Asp	Val	Lys	Met	Val	Ser	Gly	Phe	Ile	Pro	Leu	Lys
65					70					75					80
Pro	Thr	Val	Lys	Met	Leu	Glu	Arg	Ser	Asn	His	Val	Ser	Arg	Thr	Glu
				85					90					95	
Val	Ser	Ser	Asn	His	Val	Leu	Ile	Tyr	Leu	Asp	Lys	Val	Ser	Asn	Gln
			100					105					110		
Thr	Leu	Ser	Leu	Phe	Phe	Thr	Val	Leu	Gln	Asp	Val	Pro	Val	Arg	Asp
		115					120					125			
Leu	Lys	Pro	Ala	Ile	Val	Lys	Val	Tyr	Asp						
	130					135									

<210> 10
 <211> 27
 <212> PRT
 <213> Homo sapiens

Ser	Val	Ser	Tyr	Thr	Gly	Ser	Arg	Ser	Ala	Ser	Asn	Met	Ala	Ile	Val
1				5					10					15	
Asp	Val	Lys	Met	Val	Ser	Gly	Phe	Ile	Pro	Leu					
			20					25							

<210> 11
 <211> 126
 <212> PRT
 <213> Homo sapiens

Leu	Gln	Gln	Val	Ser	Leu	Pro	Glu	Leu	Pro	Gly	Glu	Tyr	Ser	Met	Lys
-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----

<212> PRT
 <213> Homo sapiens

<400> 14
 Gln Thr Ser Leu Lys Tyr Asn Ile Leu Pro Glu Lys Glu Glu Phe Pro
 1 5 10 15
 Phe Ala Leu Gly Val Gln Thr Leu Pro Gln Thr Cys Asp Glu Pro Lys
 20 25 30
 Ala His Thr Ser Phe Gln Ile Ser Leu Ser Val Ser Tyr Thr Gly Ser
 35 40 45
 Arg Ser Ala Ser Asn Met Ala Ile Val Asp Val Lys Met Val Ser Gly
 50 55 60
 Phe Ile Pro Leu Lys Pro Thr Val Lys Met Leu Glu Arg Ser Asn His
 65 70 75 80
 Val Ser Arg Thr Glu Val Ser Ser Asn His Val Leu Ile Tyr Leu Asp
 85 90 95
 Lys Val Ser Asn Gln
 100

<210> 15
 <211> 76
 <212> PRT
 <213> Homo sapiens

<400> 15
 Gln Thr Ser Leu Lys Tyr Asn Ile Leu Pro Glu Lys Glu Glu Phe Pro
 1 5 10 15
 Phe Ala Leu Gly Val Gln Thr Leu Pro Gln Thr Cys Asp Glu Pro Lys
 20 25 30
 Ala His Thr Ser Phe Gln Ile Ser Leu Ser Val Ser Tyr Thr Gly Ser
 35 40 45
 Arg Ser Ala Ser Asn Met Ala Ile Val Asp Val Lys Met Val Ser Gly
 50 55 60
 Phe Ile Pro Leu Lys Pro Thr Val Lys Met Leu Glu
 65 70 75

<210> 16
 <211> 56
 <212> PRT
 <213> Homo sapiens

<400> 16
 Gln Thr Ser Leu Lys Tyr Asn Ile Leu Pro Glu Lys Glu Glu Phe Pro
 1 5 10 15
 Phe Ala Leu Gly Val Gln Thr Leu Pro Gln Thr Cys Asp Glu Pro Lys
 20 25 30
 Ala His Thr Ser Phe Gln Ile Ser Leu Ser Val Ser Tyr Thr Gly Ser
 35 40 45
 Arg Ser Ala Ser Asn Met Ala Ile
 50 55

<210> 17
 <211> 76
 <212> PRT
 <213> Homo sapiens

<400> 17

Gln Thr Cys Asp Glu Pro Lys Ala His Thr Ser Phe Gln Ile Ser Leu
 1 5 10 15
 Ser Val Ser Tyr Thr Gly Ser Arg Ser Ala Ser Asn Met Ala Ile Val
 20 25 30
 Asp Val Lys Met Val Ser Gly Phe Ile Pro Leu Lys Pro Thr Val Lys
 35 40 45
 Met Leu Glu Arg Ser Asn His Val Ser Arg Thr Glu Val Ser Ser Asn
 50 55 60
 His Val Leu Ile Tyr Leu Asp Lys Val Ser Asn Gln
 65 70 75

<210> 18
 <211> 76
 <212> PRT
 <213> Homo sapiens

<400> 18
 Gln Thr Cys Asp Glu Pro Lys Ala His Thr Ser Phe Gln Ile Ser Leu
 1 5 10 15
 Ser Val Ser Tyr Thr Gly Ser Arg Ser Ala Ser Asn Met Ala Ile Val
 20 25 30
 Asp Val Lys Met Val Ser Gly Phe Ile Pro Leu Lys Pro Thr Val Lys
 35 40 45
 Met Leu Glu Arg Ser Asn His Val Ser Arg Thr Glu Val Ser Ser Asn
 50 55 60
 His Val Leu Ile Tyr Leu Asp Lys Val Ser Asn Gln
 65 70 75

<210> 19
 <211> 31
 <212> PRT
 <213> Homo sapiens

<400> 19
 Gln Thr Cys Asp Glu Pro Lys Ala His Thr Ser Phe Gln Ile Ser Leu
 1 5 10 15
 Ser Val Ser Tyr Thr Gly Ser Arg Ser Ala Ser Asn Met Ala Ile
 20 25 30

<210> 20
 <211> 44
 <212> PRT
 <213> Homo sapiens

<400> 20
 Lys Thr Cys Ser Pro Lys Gln Phe Ala Cys Arg Asp Gln Ile Thr Cys
 1 5 10 15
 Ile Ser Lys Gly Trp Arg Cys Asp Gly Glu Arg Asp Cys Pro Asp Gly
 20 25 30
 Ser Asp Glu Ala Pro Glu Ile Cys Pro Gln Ser Lys
 35 40

<210> 21
 <211> 86
 <212> PRT
 <213> Homo sapiens

<400> 21

Lys	Thr	Cys	Ser	Pro	Lys	Gln	Phe	Ala	Cys	Arg	Asp	Gln	Ile	Thr	Cys
1				5					10					15	
Ile	Ser	Lys	Gly	Trp	Arg	Cys	Asp	Gly	Glu	Arg	Asp	Cys	Pro	Asp	Gly
			20					25					30		
Ser	Asp	Glu	Ala	Pro	Glu	Ile	Cys	Pro	Gln	Ser	Lys	Ala	Gln	Arg	Cys
		35					40					45			
Gln	Pro	Asn	Glu	His	Asn	Cys	Leu	Gly	Thr	Glu	Leu	Cys	Val	Pro	Met
	50					55					60				
Ser	Arg	Leu	Cys	Asn	Gly	Val	Gln	Asp	Cys	Met	Asp	Gly	Ser	Asp	Glu
65					70					75					80
Gly	Pro	His	Cys	Arg	Glu										
				85											

<210> 22
 <211> 43
 <212> PRT
 <213> Homo sapiens

<400> 22

Lys	Ala	Gln	Arg	Cys	Gln	Pro	Asn	Glu	His	Asn	Cys	Leu	Gly	Thr	Glu
1				5					10					15	
Leu	Cys	Val	Pro	Met	Ser	Arg	Leu	Cys	Asn	Gly	Val	Gln	Asp	Cys	Met
			20					25					30		
Asp	Gly	Ser	Asp	Glu	Gly	Pro	His	Cys	Arg	Glu					
		35					40								

<210> 23
 <211> 42
 <212> PRT
 <213> Homo sapiens

<400> 23

Gln	Cys	Gln	Pro	Gly	Glu	Phe	Ala	Cys	Ala	Asn	Ser	Arg	Cys	Ile	Gln
1				5					10					15	
Glu	Arg	Trp	Lys	Cys	Asp	Gly	Asp	Asn	Asp	Cys	Leu	Asp	Asn	Ser	Asp
			20					25					30		
Glu	Ala	Pro	Ala	Leu	Cys	His	Gln	His	Thr						
		35					40								

<210> 24
 <211> 82
 <212> PRT
 <213> Homo sapiens

<400> 24

Gln	Cys	Gln	Pro	Gly	Glu	Phe	Ala	Cys	Ala	Asn	Ser	Arg	Cys	Ile	Gln
1				5					10					15	
Glu	Arg	Trp	Lys	Cys	Asp	Gly	Asp	Asn	Asp	Cys	Leu	Asp	Asn	Ser	Asp
			20					25					30		
Glu	Ala	Pro	Ala	Leu	Cys	His	Gln	His	Thr	Cys	Pro	Ser	Asp	Arg	Phe
		35					40					45			
Lys	Cys	Glu	Asn	Asn	Arg	Cys	Ile	Pro	Asn	Arg	Trp	Leu	Cys	Asp	Gly
	50					55					60				
Asp	Asn	Asp	Cys	Gly	Asn	Ser	Glu	Asp	Glu	Ser	Asn	Ala	Thr	Cys	Ser
65					70					75					80
Ala	Arg														

<210> 25
 <211> 122
 <212> PRT
 <213> Homo sapiens

<400> 25
 Gln Cys Gln Pro Gly Glu Phe Ala Cys Ala Asn Ser Arg Cys Ile Gln
 1 5 10 15
 Glu Arg Trp Lys Cys Asp Gly Asp Asn Asp Cys Leu Asp Asn Ser Asp
 20 25 30
 Glu Ala Pro Ala Leu Cys His Gln His Thr Cys Pro Ser Asp Arg Phe
 35 40 45
 Lys Cys Glu Asn Asn Arg Cys Ile Pro Asn Arg Trp Leu Cys Asp Gly
 50 55 60
 Asp Asn Asp Cys Gly Asn Ser Glu Asp Glu Ser Asn Ala Thr Cys Ser
 65 70 75 80
 Ala Arg Thr Cys Pro Pro Asn Gln Phe Ser Cys Ala Ser Gly Arg Cys
 85 90 95
 Ile Pro Ile Ser Trp Thr Cys Asp Leu Asp Asp Asp Cys Gly Asp Arg
 100 105 110
 Ser Asp Glu Ser Ala Ser Cys Ala Tyr Pro
 115 120

<210> 26
 <211> 161
 <212> PRT
 <213> Homo sapiens

<400> 26
 Gln Cys Gln Pro Gly Glu Phe Ala Cys Ala Asn Ser Arg Cys Ile Gln
 1 5 10 15
 Glu Arg Trp Lys Cys Asp Gly Asp Asn Asp Cys Leu Asp Asn Ser Asp
 20 25 30
 Glu Ala Pro Ala Leu Cys His Gln His Thr Cys Pro Ser Asp Arg Phe
 35 40 45
 Lys Cys Glu Asn Asn Arg Cys Ile Pro Asn Arg Trp Leu Cys Asp Gly
 50 55 60
 Asp Asn Asp Cys Gly Asn Ser Glu Asp Glu Ser Asn Ala Thr Cys Ser
 65 70 75 80
 Ala Arg Thr Cys Pro Pro Asn Gln Phe Ser Cys Ala Ser Gly Arg Cys
 85 90 95
 Ile Pro Ile Ser Trp Thr Cys Asp Leu Asp Asp Asp Cys Gly Asp Arg
 100 105 110
 Ser Asp Glu Ser Ala Ser Cys Ala Tyr Pro Thr Cys Phe Pro Leu Thr
 115 120 125
 Gln Phe Thr Cys Asn Asn Gly Arg Cys Ile Asn Ile Asn Trp Arg Cys
 130 135 140
 Asp Asn Asp Asn Asp Cys Gly Asp Asn Ser Asp Glu Ala Gly Cys Ser
 145 150 155 160
 His

<210> 27
 <211> 208
 <212> PRT
 <213> Homo sapiens

<400> 27

Gln	Cys	Gln	Pro	Gly	Glu	Phe	Ala	Cys	Ala	Asn	Ser	Arg	Cys	Ile	Gln
1				5				10						15	
Glu	Arg	Trp	Lys	Cys	Asp	Gly	Asp	Asn	Asp	Cys	Leu	Asp	Asn	Ser	Asp
			20					25					30		
Glu	Ala	Pro	Ala	Leu	Cys	His	Gln	His	Thr	Cys	Pro	Ser	Asp	Arg	Phe
		35					40					45			
Lys	Cys	Glu	Asn	Asn	Arg	Cys	Ile	Pro	Asn	Arg	Trp	Leu	Cys	Asp	Gly
	50					55					60				
Asp	Asn	Asp	Cys	Gly	Asn	Ser	Glu	Asp	Glu	Ser	Asn	Ala	Thr	Cys	Ser
65					70					75					80
Ala	Arg	Thr	Cys	Pro	Pro	Asn	Gln	Phe	Ser	Cys	Ala	Ser	Gly	Arg	Cys
				85					90					95	
Ile	Pro	Ile	Ser	Trp	Thr	Cys	Asp	Leu	Asp	Asp	Asp	Cys	Gly	Asp	Arg
			100					105					110		
Ser	Asp	Glu	Ser	Ala	Ser	Cys	Ala	Tyr	Pro	Thr	Cys	Phe	Pro	Leu	Thr
		115					120					125			
Gln	Phe	Thr	Cys	Asn	Asn	Gly	Arg	Cys	Ile	Asn	Ile	Asn	Trp	Arg	Cys
	130					135					140				
Asp	Asn	Asp	Asn	Asp	Cys	Gly	Asp	Asn	Ser	Asp	Glu	Ala	Gly	Cys	Ser
145					150					155					160
His	Ser	Cys	Ser	Ser	Thr	Gln	Phe	Lys	Cys	Asn	Ser	Gly	Arg	Cys	Ile
				165					170					175	
Pro	Glu	His	Trp	Thr	Cys	Asp	Gly	Asp	Asn	Asp	Cys	Gly	Asp	Tyr	Ser
			180					185					190		
Asp	Glu	Thr	His	Ala	Asn	Cys	Thr	Asn	Gln	Ala	Thr	Arg	Pro	Pro	Gly
		195					200					205			

<210> 28
 <211> 150
 <212> PRT
 <213> Homo sapiens

Gln	Cys	Gln	Pro	Gly	Glu	Phe	Ala	Cys	Ala	Asn	Ser	Arg	Cys	Ile	Gln
1				5				10						15	
Glu	Arg	Trp	Lys	Cys	Asp	Gly	Asp	Asn	Asp	Cys	Leu	Asp	Asn	Ser	Asp
			20					25					30		
Glu	Ala	Pro	Ala	Leu	Cys	His	Gln	His	Thr	Cys	Pro	Ser	Asp	Arg	Phe
		35					40					45			
Lys	Cys	Glu	Asn	Asn	Arg	Cys	Ile	Pro	Asn	Arg	Trp	Leu	Cys	Asp	Gly
	50					55					60				
Asp	Asn	Asp	Cys	Gly	Asn	Ser	Glu	Asp	Glu	Ser	Asn	Ala	Thr	Cys	Ser
65					70					75					80
Ala	Arg	Thr	Cys	Pro	Pro	Asn	Gln	Phe	Ser	Cys	Ala	Ser	Gly	Arg	Cys
				85					90					95	
Ile	Pro	Ile	Ser	Trp	Thr	Cys	Asp	Leu	Asp	Asp	Asp	Cys	Gly	Asp	Arg
			100					105					110		
Ser	Asp	Glu	Ser	Ala	Ser	Cys	Ala	Tyr	Pro	Thr	Cys	Phe	Pro	Leu	Thr
		115					120					125			
Gln	Phe	Thr	Cys	Asn	Asn	Gly	Arg	Cys	Ile	Asn	Ile	Asn	Trp	Arg	Cys
	130					135					140				
Asp	Asn	Asp	Asn	Asp	Cys										
145					150										

<210> 29
 <211> 231
 <212> PRT
 <213> Homo sapiens

<400> 29
 Gln Cys Gln Pro Gly Glu Phe Ala Cys Ala Asn Ser Arg Cys Ile Gln
 1 5 10 15
 Glu Arg Trp Lys Cys Asp Gly Asp Asn Asp Cys Leu Asp Asn Ser Asp
 20 25 30
 Glu Ala Pro Ala Leu Cys His Gln His Thr Cys Pro Ser Asp Arg Phe
 35 40 45
 Lys Cys Glu Asn Asn Arg Cys Ile Pro Asn Arg Trp Leu Cys Asp Gly
 50 55 60
 Asp Asn Asp Cys Gly Asn Ser Glu Asp Glu Ser Asn Ala Thr Cys Ser
 65 70 75 80
 Ala Arg Thr Cys Pro Pro Asn Gln Phe Ser Cys Ala Ser Gly Arg Cys
 85 90 95
 Ile Pro Ile Ser Trp Thr Cys Asp Leu Asp Asp Asp Cys Gly Asp Arg
 100 105 110
 Ser Asp Glu Ser Ala Ser Cys Ala Tyr Pro Thr Cys Phe Pro Leu Thr
 115 120 125
 Gln Phe Thr Cys Asn Asn Gly Arg Cys Ile Asn Ile Asn Trp Arg Cys
 130 135 140
 Asp Asn Asp Asn Asp Cys Gly Asp Asn Ser Asp Glu Ala Gly Cys Ser
 145 150 155 160
 His Ser Cys Ser Ser Thr Gln Phe Lys Cys Asn Ser Gly Arg Cys Ile
 165 170 175
 Pro Glu His Trp Thr Cys Asp Gly Asp Asn Asp Cys Gly Asp Tyr Ser
 180 185 190
 Asp Glu Thr His Ala Asn Cys Thr Asn Gln Ala Thr Arg Pro Pro Gly
 195 200 205
 Gly Cys His Thr Asp Glu Phe Gln Cys Arg Leu Asp Gly Leu Cys Ile
 210 215 220
 Pro Leu Arg Trp Arg Cys Asp
 225 230

<210> 30
 <211> 40
 <212> PRT
 <213> Homo sapiens

<400> 30
 Cys Pro Ser Asp Arg Phe Lys Cys Glu Asn Asn Arg Cys Ile Pro Asn
 1 5 10 15
 Arg Trp Leu Cys Asp Gly Asp Asn Asp Cys Gly Asn Ser Glu Asp Glu
 20 25 30
 Ser Asn Ala Thr Cys Ser Ala Arg
 35 40

<210> 31
 <211> 80
 <212> PRT
 <213> Homo sapiens

<400> 31
 Cys Pro Ser Asp Arg Phe Lys Cys Glu Asn Asn Arg Cys Ile Pro Asn
 1 5 10 15
 Arg Trp Leu Cys Asp Gly Asp Asn Asp Cys Gly Asn Ser Glu Asp Glu
 20 25 30
 Ser Asn Ala Thr Cys Ser Ala Arg Thr Cys Pro Pro Asn Gln Phe Ser
 35 40 45
 Cys Ala Ser Gly Arg Cys Ile Pro Ile Ser Trp Thr Cys Asp Leu Asp

50 55 60
 Asp Asp Cys Gly Asp Arg Ser Asp Glu Ser Ala Ser Cys Ala Tyr Pro
 65 70 75 80

<210> 32
 <211> 119
 <212> PRT
 <213> Homo sapiens

<400> 32
 Cys Pro Ser Asp Arg Phe Lys Cys Glu Asn Asn Arg Cys Ile Pro Asn
 1 5 10 15
 Arg Trp Leu Cys Asp Gly Asp Asn Asp Cys Gly Asn Ser Glu Asp Glu
 20 25 30
 Ser Asn Ala Thr Cys Ser Ala Arg Thr Cys Pro Pro Asn Gln Phe Ser
 35 40 45
 Cys Ala Ser Gly Arg Cys Ile Pro Ile Ser Trp Thr Cys Asp Leu Asp
 50 55 60
 Asp Asp Cys Gly Asp Arg Ser Asp Glu Ser Ala Ser Cys Ala Tyr Pro
 65 70 75 80
 Thr Cys Phe Pro Leu Thr Gln Phe Thr Cys Asn Asn Gly Arg Cys Ile
 85 90 95
 Asn Ile Asn Trp Arg Cys Asp Asn Asp Asn Cys Gly Asp Asn Ser
 100 105 110
 Asp Glu Ala Gly Cys Ser His
 115

<210> 33
 <211> 166
 <212> PRT
 <213> Homo sapiens

<400> 33
 Cys Pro Ser Asp Arg Phe Lys Cys Glu Asn Asn Arg Cys Ile Pro Asn
 1 5 10 15
 Arg Trp Leu Cys Asp Gly Asp Asn Asp Cys Gly Asn Ser Glu Asp Glu
 20 25 30
 Ser Asn Ala Thr Cys Ser Ala Arg Thr Cys Pro Pro Asn Gln Phe Ser
 35 40 45
 Cys Ala Ser Gly Arg Cys Ile Pro Ile Ser Trp Thr Cys Asp Leu Asp
 50 55 60
 Asp Asp Cys Gly Asp Arg Ser Asp Glu Ser Ala Ser Cys Ala Tyr Pro
 65 70 75 80
 Thr Cys Phe Pro Leu Thr Gln Phe Thr Cys Asn Asn Gly Arg Cys Ile
 85 90 95
 Asn Ile Asn Trp Arg Cys Asp Asn Asp Asn Cys Gly Asp Asn Ser
 100 105 110
 Asp Glu Ala Gly Cys Ser His Ser Cys Ser Ser Thr Gln Phe Lys Cys
 115 120 125
 Asn Ser Gly Arg Cys Ile Pro Glu His Trp Thr Cys Asp Gly Asp Asn
 130 135 140
 Asp Cys Gly Asp Tyr Ser Asp Glu Thr His Ala Asn Cys Thr Asn Gln
 145 150 155 160
 Ala Thr Arg Pro Pro Gly
 165

<210> 34
 <211> 108

<212> PRT
<213> Homo sapiens

<400> 34
Cys Pro Ser Asp Arg Phe Lys Cys Glu Asn Asn Arg Cys Ile Pro Asn
1 5 10 15
Arg Trp Leu Cys Asp Gly Asp Asn Asp Cys Gly Asn Ser Glu Asp Glu
20 25 30
Ser Asn Ala Thr Cys Ser Ala Arg Thr Cys Pro Pro Asn Gln Phe Ser
35 40 45
Cys Ala Ser Gly Arg Cys Ile Pro Ile Ser Trp Thr Cys Asp Leu Asp
50 55 60
Asp Asp Cys Gly Asp Arg Ser Asp Glu Ser Ala Ser Cys Ala Tyr Pro
65 70 75 80
Thr Cys Phe Pro Leu Thr Gln Phe Thr Cys Asn Asn Gly Arg Cys Ile
85 90 95
Asn Ile Asn Trp Arg Cys Asp Asn Asp Asn Asp Cys
100 105

<210> 35
<211> 289
<212> PRT
<213> Homo sapiens

<400> 35
Cys Pro Ser Asp Arg Phe Lys Cys Glu Asn Asn Arg Cys Ile Pro Asn
1 5 10 15
Arg Trp Leu Cys Asp Gly Asp Asn Asp Cys Gly Asn Ser Glu Asp Glu
20 25 30
Ser Asn Ala Thr Cys Ser Ala Arg Thr Cys Pro Pro Asn Gln Phe Ser
35 40 45
Cys Ala Ser Gly Arg Cys Ile Pro Ile Ser Trp Thr Cys Asp Leu Asp
50 55 60
Asp Asp Cys Gly Asp Arg Ser Asp Glu Ser Ala Ser Cys Ala Tyr Pro
65 70 75 80
Thr Cys Phe Pro Leu Thr Gln Phe Thr Cys Asn Asn Gly Arg Cys Ile
85 90 95
Asn Ile Asn Trp Arg Cys Asp Asn Asp Asn Asp Cys Gly Asp Asn Ser
100 105 110
Asp Glu Ala Gly Cys Ser His Ser Cys Ser Ser Thr Gln Phe Lys Cys
115 120 125
Asn Ser Gly Arg Cys Ile Pro Glu His Trp Thr Cys Asp Gly Asp Asn
130 135 140
Asp Cys Gly Asp Tyr Ser Asp Glu Thr His Ala Asn Cys Thr Asn Gln
145 150 155 160
Ala Thr Arg Pro Pro Gly Gly Cys His Thr Asp Glu Phe Gln Cys Arg
165 170 175
Leu Asp Gly Leu Cys Ile Pro Leu Arg Trp Arg Cys Asp Gly Asp Thr
180 185 190
Asp Cys Met Asp Ser Ser Asp Glu Lys Ser Cys Glu Gly Val Thr His
195 200 205
Val Cys Asp Pro Ser Val Lys Phe Gly Cys Lys Asp Ser Ala Arg Cys
210 215 220
Ile Ser Lys Ala Trp Val Cys Asp Gly Asp Asn Asp Cys Glu Asp Asn
225 230 235 240
Ser Asp Glu Glu Asn Cys Glu Ser Leu Ala Cys Arg Pro Pro Ser His
245 250 255

Pro Cys Ala Asn Asn Thr Ser Val Cys Leu Pro Pro Asp Lys Leu Cys
 260 265 270
 Asp Gly Asn Asp Asp Cys Gly Asp Gly Ser Asp Glu Gly Glu Leu Cys
 275 280 285
 Asp

<210> 36
 <211> 40
 <212> PRT
 <213> Homo sapiens

<400> 36
 Thr Cys Pro Pro Asn Gln Phe Ser Cys Ala Ser Gly Arg Cys Ile Pro
 1 5 10 15
 Ile Ser Trp Thr Cys Asp Leu Asp Asp Asp Cys Gly Asp Arg Ser Asp
 20 25 30
 Glu Ser Ala Ser Cys Ala Tyr Pro
 35 40

<210> 37
 <211> 79
 <212> PRT
 <213> Homo sapiens

<400> 37
 Thr Cys Pro Pro Asn Gln Phe Ser Cys Ala Ser Gly Arg Cys Ile Pro
 1 5 10 15
 Ile Ser Trp Thr Cys Asp Leu Asp Asp Asp Cys Gly Asp Arg Ser Asp
 20 25 30
 Glu Ser Ala Ser Cys Ala Tyr Pro Thr Cys Phe Pro Leu Thr Gln Phe
 35 40 45
 Thr Cys Asn Asn Gly Arg Cys Ile Asn Ile Asn Trp Arg Cys Asp Asn
 50 55 60
 Asp Asn Asp Cys Gly Asp Asn Ser Asp Glu Ala Gly Cys Ser His
 65 70 75

<210> 38
 <211> 126
 <212> PRT
 <213> Homo sapiens

<400> 38
 Thr Cys Pro Pro Asn Gln Phe Ser Cys Ala Ser Gly Arg Cys Ile Pro
 1 5 10 15
 Ile Ser Trp Thr Cys Asp Leu Asp Asp Asp Cys Gly Asp Arg Ser Asp
 20 25 30
 Glu Ser Ala Ser Cys Ala Tyr Pro Thr Cys Phe Pro Leu Thr Gln Phe
 35 40 45
 Thr Cys Asn Asn Gly Arg Cys Ile Asn Ile Asn Trp Arg Cys Asp Asn
 50 55 60
 Asp Asn Asp Cys Gly Asp Asn Ser Asp Glu Ala Gly Cys Ser His Ser
 65 70 75 80
 Cys Ser Ser Thr Gln Phe Lys Cys Asn Ser Gly Arg Cys Ile Pro Glu
 85 90 95
 His Trp Thr Cys Asp Gly Asp Asn Asp Cys Gly Asp Tyr Ser Asp Glu
 100 105 110
 Thr His Ala Asn Cys Thr Asn Gln Ala Thr Arg Pro Pro Gly

115	120	125
<210> 39		
<211> 68		
<212> PRT		
<213> Homo sapiens		
<400> 39		
Thr Cys Pro Pro Asn Gln Phe Ser Cys Ala Ser Gly Arg Cys Ile Pro		
1 5 10 15		
Ile Ser Trp Thr Cys Asp Leu Asp Asp Asp Cys Gly Asp Arg Ser Asp		
20 25 30		
Glu Ser Ala Ser Cys Ala Tyr Pro Thr Cys Phe Pro Leu Thr Gln Phe		
35 40 45		
Thr Cys Asn Asn Gly Arg Cys Ile Asn Ile Asn Trp Arg Cys Asp Asn		
50 55 60		
Asp Asn Asp Cys		
65		
<210> 40		
<211> 248		
<212> PRT		
<213> Homo sapiens		
<400> 40		
Cys Pro Pro Asn Gln Phe Ser Cys Ala Ser Gly Arg Cys Ile Pro Ile		
1 5 10 15		
Ser Trp Thr Cys Asp Leu Asp Asp Asp Cys Gly Asp Arg Ser Asp Glu		
20 25 30		
Ser Ala Ser Cys Ala Tyr Pro Thr Cys Phe Pro Leu Thr Gln Phe Thr		
35 40 45		
Cys Asn Asn Gly Arg Cys Ile Asn Ile Asn Trp Arg Cys Asp Asn Asp		
50 55 60		
Asn Asp Cys Gly Asp Asn Ser Asp Glu Ala Gly Cys Ser His Ser Cys		
65 70 75 80		
Ser Ser Thr Gln Phe Lys Cys Asn Ser Gly Arg Cys Ile Pro Glu His		
85 90 95		
Trp Thr Cys Asp Gly Asp Asn Asp Cys Gly Asp Tyr Ser Asp Glu Thr		
100 105 110		
His Ala Asn Cys Thr Asn Gln Ala Thr Arg Pro Pro Gly Gly Cys His		
115 120 125		
Thr Asp Glu Phe Gln Cys Arg Leu Asp Gly Leu Cys Ile Pro Leu Arg		
130 135 140		
Trp Arg Cys Asp Gly Asp Thr Asp Cys Met Asp Ser Ser Asp Glu Lys		
145 150 155 160		
Ser Cys Glu Gly Val Thr His Val Cys Asp Pro Ser Val Lys Phe Gly		
165 170 175		
Cys Lys Asp Ser Ala Arg Cys Ile Ser Lys Ala Trp Val Cys Asp Gly		
180 185 190		
Asp Asn Asp Cys Glu Asp Asn Ser Asp Glu Glu Asn Cys Glu Ser Leu		
195 200 205		
Ala Cys Arg Pro Pro Ser His Pro Cys Ala Asn Asn Thr Ser Val Cys		
210 215 220		
Leu Pro Pro Asp Lys Leu Cys Asp Gly Asn Asp Asp Cys Gly Asp Gly		
225 230 235 240		
Ser Asp Glu Gly Glu Leu Cys Asp		
245		

<210> 41
 <211> 39
 <212> PRT
 <213> Homo sapiens

<400> 41
 Thr Cys Phe Pro Leu Thr Gln Phe Thr Cys Asn Asn Gly Arg Cys Ile
 1 5 10 15
 Asn Ile Asn Trp Arg Cys Asp Asn Asp Asn Asp Cys Gly Asp Asn Ser
 20 25 30
 Asp Glu Ala Gly Cys Ser His
 35

<210> 42
 <211> 86
 <212> PRT
 <213> Homo sapiens

<400> 42
 Thr Cys Phe Pro Leu Thr Gln Phe Thr Cys Asn Asn Gly Arg Cys Ile
 1 5 10 15
 Asn Ile Asn Trp Arg Cys Asp Asn Asp Asn Asp Cys Gly Asp Asn Ser
 20 25 30
 Asp Glu Ala Gly Cys Ser His Ser Cys Ser Ser Thr Gln Phe Lys Cys
 35 40 45
 Asn Ser Gly Arg Cys Ile Pro Glu His Trp Thr Cys Asp Gly Asp Asn
 50 55 60
 Asp Cys Gly Asp Tyr Ser Asp Glu Thr His Ala Asn Cys Thr Asn Gln
 65 70 75 80
 Ala Thr Arg Pro Pro Gly
 85

<210> 43
 <211> 169
 <212> PRT
 <213> Homo sapiens

<400> 43
 Thr Cys Phe Pro Leu Thr Gln Phe Thr Cys Asn Asn Gly Arg Cys Ile
 1 5 10 15
 Asn Ile Asn Trp Arg Cys Asp Asn Asp Asn Asp Cys Gly Asp Asn Ser
 20 25 30
 Asp Glu Ala Gly Cys Ser His Ser Cys Ser Ser Thr Gln Phe Lys Cys
 35 40 45
 Asn Ser Gly Arg Cys Ile Pro Glu His Trp Thr Cys Asp Gly Asp Asn
 50 55 60
 Asp Cys Gly Asp Tyr Ser Asp Glu Thr His Ala Asn Cys Thr Asn Gln
 65 70 75 80
 Ala Thr Arg Pro Pro Gly Gly Cys His Thr Asp Glu Phe Gln Cys Arg
 85 90 95
 Leu Asp Gly Leu Cys Ile Pro Leu Arg Trp Arg Cys Asp Gly Asp Thr
 100 105 110
 Asp Cys Met Asp Ser Ser Asp Glu Lys Ser Cys Glu Gly Val Thr His
 115 120 125
 Val Cys Asp Pro Ser Val Lys Phe Gly Cys Lys Asp Ser Ala Arg Cys
 130 135 140
 Ile Ser Lys Ala Trp Val Cys Asp Gly Asp Asn Asp Cys Glu Asp Asn
 145 150 155 160

Ser Asp Glu Glu Asn Cys Glu Ser Leu
165

<210> 44
<211> 209
<212> PRT
<213> Homo sapiens

<400> 44
Thr Cys Phe Pro Leu Thr Gln Phe Thr Cys Asn Asn Gly Arg Cys Ile
1 5 10 15
Asn Ile Asn Trp Arg Cys Asp Asn Asp Asn Asp Cys Gly Asp Asn Ser
20 25 30
Asp Glu Ala Gly Cys Ser His Ser Cys Ser Ser Thr Gln Phe Lys Cys
35 40 45
Asn Ser Gly Arg Cys Ile Pro Glu His Trp Thr Cys Asp Gly Asp Asn
50 55 60
Asp Cys Gly Asp Tyr Ser Asp Glu Thr His Ala Asn Cys Thr Asn Gln
65 70 75 80
Ala Thr Arg Pro Pro Gly Gly Cys His Thr Asp Glu Phe Gln Cys Arg
85 90 95
Leu Asp Gly Leu Cys Ile Pro Leu Arg Trp Arg Cys Asp Gly Asp Thr
100 105 110
Asp Cys Met Asp Ser Ser Asp Glu Lys Ser Cys Glu Gly Val Thr His
115 120 125
Val Cys Asp Pro Ser Val Lys Phe Gly Cys Lys Asp Ser Ala Arg Cys
130 135 140
Ile Ser Lys Ala Trp Val Cys Asp Gly Asp Asn Asp Cys Glu Asp Asn
145 150 155 160
Ser Asp Glu Glu Asn Cys Glu Ser Leu Ala Cys Arg Pro Pro Ser His
165 170 175
Pro Cys Ala Asn Asn Thr Ser Val Cys Leu Pro Pro Asp Lys Leu Cys
180 185 190
Asp Gly Asn Asp Asp Cys Gly Asp Gly Ser Asp Glu Gly Glu Leu Cys
195 200 205
Asp

<210> 45
<211> 47
<212> PRT
<213> Homo sapiens

<400> 45
Ser Cys Ser Ser Thr Gln Phe Lys Cys Asn Ser Gly Arg Cys Ile Pro
1 5 10 15
Glu His Trp Thr Cys Asp Gly Asp Asn Asp Cys Gly Asp Tyr Ser Asp
20 25 30
Glu Thr His Ala Asn Cys Thr Asn Gln Ala Thr Arg Pro Pro Gly
35 40 45

<210> 46
<211> 89
<212> PRT
<213> Homo sapiens

<400> 46
Ser Cys Ser Ser Thr Gln Phe Lys Cys Asn Ser Gly Arg Cys Ile Pro

1		5		10		15									
Glu	His	Trp	Thr	Cys	Asp	Gly	Asp	Asn	Asp	Cys	Gly	Asp	Tyr	Ser	Asp
		20						25					30		
Glu	Thr	His	Ala	Asn	Cys	Thr	Asn	Gln	Ala	Thr	Arg	Pro	Pro	Gly	Gly
		35						40					45		
Cys	His	Thr	Asp	Glu	Phe	Gln	Cys	Arg	Leu	Asp	Gly	Leu	Cys	Ile	Pro
		50				55					60				
Leu	Arg	Trp	Arg	Cys	Asp	Gly	Asp	Thr	Asp	Cys	Met	Asp	Ser	Ser	Asp
65					70					75					80
Glu	Lys	Ser	Cys	Glu	Gly	Val	Thr	His							
				85											

<210> 47
 <211> 170
 <212> PRT
 <213> Homo sapiens

	<400> 47														
Ser	Cys	Ser	Ser	Thr	Gln	Phe	Lys	Cys	Asn	Ser	Gly	Arg	Cys	Ile	Pro
1		5						10					15		
Glu	His	Trp	Thr	Cys	Asp	Gly	Asp	Asn	Asp	Cys	Gly	Asp	Tyr	Ser	Asp
		20						25					30		
Glu	Thr	His	Ala	Asn	Cys	Thr	Asn	Gln	Ala	Thr	Arg	Pro	Pro	Gly	Gly
		35						40					45		
Cys	His	Thr	Asp	Glu	Phe	Gln	Cys	Arg	Leu	Asp	Gly	Leu	Cys	Ile	Pro
		50				55					60				
Leu	Arg	Trp	Arg	Cys	Asp	Gly	Asp	Thr	Asp	Cys	Met	Asp	Ser	Ser	Asp
65					70					75					80
Glu	Lys	Ser	Cys	Glu	Gly	Val	Thr	His	Val	Cys	Asp	Pro	Ser	Val	Lys
				85					90					95	
Phe	Gly	Cys	Lys	Asp	Ser	Ala	Arg	Cys	Ile	Ser	Lys	Ala	Trp	Val	Cys
			100					105					110		
Asp	Gly	Asp	Asn	Asp	Cys	Glu	Asp	Asn	Ser	Asp	Glu	Glu	Asn	Cys	Glu
		115					120					125			
Ser	Leu	Ala	Cys	Arg	Pro	Pro	Ser	His	Pro	Cys	Ala	Asn	Asn	Thr	Ser
		130				135					140				
Val	Cys	Leu	Pro	Pro	Asp	Lys	Leu	Cys	Asp	Gly	Asn	Asp	Asp	Cys	Gly
145					150					155					160
Asp	Gly	Ser	Asp	Glu	Gly	Glu	Leu	Cys	Asp						
				165					170						

<210> 48
 <211> 42
 <212> PRT
 <213> Homo sapiens

	<400> 48														
Gly	Cys	His	Thr	Asp	Glu	Phe	Gln	Cys	Arg	Leu	Asp	Gly	Leu	Cys	Ile
1		5						10					15		
Pro	Leu	Arg	Trp	Arg	Cys	Asp	Gly	Asp	Thr	Asp	Cys	Met	Asp	Ser	Ser
		20						25					30		
Asp	Glu	Lys	Ser	Cys	Glu	Gly	Val	Thr	His						
		35					40								

<210> 49
 <211> 83
 <212> PRT
 <213> Homo sapiens

<400> 49
 Gly Cys His Thr Asp Glu Phe Gln Cys Arg Leu Asp Gly Leu Cys Ile
 1 5 10 15
 Pro Leu Arg Trp Arg Cys Asp Gly Asp Thr Asp Cys Met Asp Ser Ser
 20 25 30
 Asp Glu Lys Ser Cys Glu Gly Val Thr His Val Cys Asp Pro Ser Val
 35 40 45
 Lys Phe Gly Cys Lys Asp Ser Ala Arg Cys Ile Ser Lys Ala Trp Val
 50 55 60
 Cys Asp Gly Asp Asn Asp Cys Glu Asp Asn Ser Asp Glu Glu Asn Cys
 65 70 75 80
 Glu Ser Leu

<210> 50
 <211> 123
 <212> PRT
 <213> Homo sapiens

<400> 50
 Gly Cys His Thr Asp Glu Phe Gln Cys Arg Leu Asp Gly Leu Cys Ile
 1 5 10 15
 Pro Leu Arg Trp Arg Cys Asp Gly Asp Thr Asp Cys Met Asp Ser Ser
 20 25 30
 Asp Glu Lys Ser Cys Glu Gly Val Thr His Val Cys Asp Pro Ser Val
 35 40 45
 Lys Phe Gly Cys Lys Asp Ser Ala Arg Cys Ile Ser Lys Ala Trp Val
 50 55 60
 Cys Asp Gly Asp Asn Asp Cys Glu Asp Asn Ser Asp Glu Glu Asn Cys
 65 70 75 80
 Glu Ser Leu Ala Cys Arg Pro Pro Ser His Pro Cys Ala Asn Asn Thr
 85 90 95
 Ser Val Cys Leu Pro Pro Asp Lys Leu Cys Asp Gly Asn Asp Asp Cys
 100 105 110
 Gly Asp Gly Ser Asp Glu Gly Glu Leu Cys Asp
 115 120

<210> 51
 <211> 41
 <212> PRT
 <213> Homo sapiens

<400> 51
 Val Cys Asp Pro Ser Val Lys Phe Gly Cys Lys Asp Ser Ala Arg Cys
 1 5 10 15
 Ile Ser Lys Ala Trp Val Cys Asp Gly Asp Asn Asp Cys Glu Asp Asn
 20 25 30
 Ser Asp Glu Glu Asn Cys Glu Ser Leu
 35 40

<210> 52
 <211> 81
 <212> PRT
 <213> Homo sapiens

<400> 52
 Val Cys Asp Pro Ser Val Lys Phe Gly Cys Lys Asp Ser Ala Arg Cys
 1 5 10 15

Ile	Ser	Lys	Ala	Trp	Val	Cys	Asp	Gly	Asp	Asn	Asp	Cys	Glu	Asp	Asn
			20					25					30		
Ser	Asp	Glu	Glu	Asn	Cys	Glu	Ser	Leu	Ala	Cys	Arg	Pro	Pro	Ser	His
		35					40					45			
Pro	Cys	Ala	Asn	Asn	Thr	Ser	Val	Cys	Leu	Pro	Pro	Asp	Lys	Leu	Cys
	50					55					60				
Asp	Gly	Asn	Asp	Asp	Cys	Gly	Asp	Gly	Ser	Asp	Glu	Gly	Glu	Leu	Cys
65					70					75					80
Asp															

<210> 53
 <211> 40
 <212> PRT
 <213> Homo sapiens

Ala	Cys	Arg	Pro	Pro	Ser	His	Pro	Cys	Ala	Asn	Asn	Thr	Ser	Val	Cys
1				5					10					15	
Leu	Pro	Pro	Asp	Lys	Leu	Cys	Asp	Gly	Asn	Asp	Asp	Cys	Gly	Asp	Gly
			20					25					30		
Ser	Asp	Glu	Gly	Glu	Leu	Cys	Asp								
		35					40								

<210> 54
 <211> 10
 <212> PRT
 <213> Homo sapiens

Ser	Gly	Phe	Ser	Leu	Gly	Ser	Asp	Gly	Lys						
1				5					10						

<210> 55
 <211> 10
 <212> PRT
 <213> Homo sapiens

Gly	Ile	Ala	Leu	Asp	Pro	Ala	Met	Gly	Lys						
1				5					10						

<210> 56
 <211> 10
 <212> PRT
 <213> Homo sapiens

Gly	Gly	Ala	Leu	His	Ile	Tyr	His	Gln	Arg						
1				5					10						

<210> 57
 <211> 11
 <212> PRT
 <213> Homo sapiens

<400> 57

Val Phe Phe Thr Asp Tyr Gly Gln Ile Pro Lys
1 5 10

Phe

Thr

Tyr

Gln

Pro

SECRET